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Result
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length: 2000000000
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geneseqp1990s:*
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ADL70790
AAU02980
ABR82321
ADT08174
ABM83778
ABM83783
           ADH89360
ADK70589
ADL704413
ADL24413
ADU17885
ADT55264
ADT75914
ADT789149
ADT78149
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Biocceleration Ltd.
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100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
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Ad170731	Adk15868	Adh89359	Adh34559	Adf74800	Adf74796	Add45282	Abp72819	Aay50717	Aaw54354	Aar66492	Aar12499	Aap70384	Aee47501 Human tra	Ady57103	Ady57102	Ady54149	Adt55066	Adu00630	Adu17468	Adu17611	Adt98423
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ALI GNMENTS

RESULT 1

ADG44071

ADG440 New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization. antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. Human CDT peptide fragment ADG46075 standard; peptide; WPI; 2004-073743/08. Althaus 05-JUL-2002; 19-MAY-2003; 2003EP-00011334. 07-JAN-2004. EP1378521-A1. Homo sapiens. 25-MAR-2004 (first entry) ADG46075; (DADE-) DADE BEHRING MARBURG GMBH 2002DE-01030550 #1. 17 A A

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSWGKEDLIWELL, TTEDSIAKIMNGRADAMSLDGGF, SKLSWGSGIMNSENN A YEKYLGEEYVRAV (ADG46075-ADG46079) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for

and

Claim 4; SEQ ID NO 1; 21pp; German.

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Matches 17
                                   29-AUG-2002;
17-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; Antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted protein SEQ ID
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12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ch 100.0%; Score 86; DB 8; I l Similarity 100.0%; Pred. No. 3.4e-07; 17; Conservative 0; Mismatches 0;
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                                   2002US-040666P
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2002US-0410948P
2002US-0410949P
2002US-0410959P
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2002US-0410959P
2002US-0410959P
2002US-0410959P
2002US-0410960P
2002US-0410960P
2002US-0410961P
2002US-0410961P
2002US-0411019P
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2002US-0411019P
2002US-0411019P
2002US-0411023P
2002US-0411023P
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2002US-0406642P.
2002US-0406646P.
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2002US-0406655P.
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2002US-0406616P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunosuppressive; Antibacterial; Virucide; human secreted protein.
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12 MAY 2003
02 MAY 2003
02 MAY 2003
19 JUL 2003
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11 JUL 2003
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17-SEP-2002;
17-SEP-2002;
                                                                                                                                                                      The present invention relates to an isolated nucleic acid molecule encoding a polypeptide which is believed to be cytostatic, antiinflammatory, immunosuppressive, antibacterial and virucidal. The composition and methods are useful for diagnosing, preventing and treating diseases such as proliferative (e.g. cancer), inflammatory, immune, metabolic, genetic, bacterial and viral diseases. The present sequence represents a human secreted protein. The present sequence is available on WIPOWEB and is not in the specification.
                                                                                                                                                                                                                                                                                                                                          New nucleic acid molecule for diagnosing, preventing or treating diseases such as proliferative (e.g. cancer), inflammatory, immune, metabolic, genetic, bacterial and viral diseases.
                                                                                                   Sequence
                                                                                                                               Revised record corrected
                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 2377; 428pp; English.
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Pierce K, Wa
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17-SEP-2002;
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                                                       Local
1 VVARSMGGKEDLIWELL 17
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17; Conser
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k RF, Huang |
, Wang Y, W
                                       100.0%;
llarity 100.0%;
Conservative 0,
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2003US-0493341P.
2003US-0493370P.
2003US-0493573P.
2003US-0493577P.
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2002US-0411055P
2002US-0411055P
2002US-0411101P
2002US-0411101P
2002US-0411101P
2002US-0463708P
2003US-0463716P
2003US-046373P
2003US-0467201P
2003US-0467201P
2003US-0467201P
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2003US-0472430P
2003US-0472430P
2003US-0476649P
2003US-0485218P
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2002US-0411045P.
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Wong JGP,
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                                                                                                                                                01-DEC-2005 : Sequence description line
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Kothakota S, Haishan L,
IGP, Wu G, Zhang H;
                                        Score 86; DB
Pred. No. 5.1
); Mismatches
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                                                       5.1e-06;
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L. Linnemann '
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124

VVARSMGGKEDLIWELL 140

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RESULT 4
ADL70790
ID ADL7
 ADL70790 standard; protein; 371 AA
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ADH89432
ID ADH894
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New fusion protein comprising a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region, useful functions a composition for treating e.g., septic shock, neoplasm or
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30-NUV-2001; 2001US-0334059P.
30-AUG-2002; 2002US-00231494.
30-AUG-2002; 2002US-0406977P.
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                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                               invention.
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183
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17; Conserv
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100.0%; Pred.
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Novel library containing several fusion proteins each of which comprises first transferrin polypeptide fused to at least one second peptide, useful for screening for transferrin fusion protein having the particular
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N-PSDB; ADL70789.
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10-MAR-2003; 2003US-003B4060
09-JUL-2003; 2003US-0485404P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunosuppressive; Haemostatic; Antiallergic; Antiasthmatic; Dermatological; Antiinflammatory; Antibacterial; Vasotropic; Nephrotropic; Neuroprotective; Cytostatic; Cerebroprotective; Vulnerary; Antiparkinsonian; Nootropic; Cardiant; Antianaemic; Antiatherosclerotic; Thrombolytic; Anticoagulant; Gastrointestinal; Respiratory; transferrin;
                                                                                                                                                                                                                                                                                                                                                                                                          Prior CP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-2003; 2003WO-US026779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        If; transferrin fusion protein; If fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transferrin N-domain protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL70790;
                                                                                                                                                                                                                                                                                                                                                                                                                           (BIOR-) BIOREXIS PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                         Turner AJ,
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                                                                                                                                                                                                                                                                                                                                                                                                         Sadeghi
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Example 13; SEQ ID NO 63; 243pp; English.

CC disorders or diseases of endocrine system, nervous system, inmune system, crespiratory system, cardiovascular system, diseases and/or disorders or diseases of endocrine system, diseases and/or disorders (CC provention and/or treatment of autoimmune disorders; diseases and disorders of hasmatopoietic cells (e.g., leukopenia, neutropenia, anaemia and thrombocytopenia); allergic reactions such as altergic asthma, cc anaphylaxis, IgE-mediated allergic reactions such as altergic asthma, rhinitis and ezema; inflammatory conditions e.g., inflammation associated with cc infection (e.g., septic shock, sepsis), ischaemia-reperfusion injury, cnephritis, Crohn's disease, multiple sclerosis, respiratory disorders (multiple sclerosis, respiratory disorders (multiple sclerosis, stroke, traumatic brain injury, neurodegenerative disorders such as Parkinson's disease, Alzheimer's disease), etc. The cusion protein is also useful as an adjuvant to enhance antibacterial or antifungal immune responses, antiparasitic immune responses, etc. The cusion protein is also useful for treating monoclonal gammopathy of cadult respiratory distress syndrome, for strumlating wound repair, for coreventing or treating infections of joints, bones, skin, etc. The fusion protein is also useful for treating or preventing thrombosytomas, it has a preventing thrombosis, myocardial construing ship career, thrombosytopenia, sickle cell anaemia, myocardial construing allows, etc. The fusion protein is also useful for treating or preventing thrombosis, myocardial construing peptide sequences uses the N-domain of If incorporated dispite the pull protein, and the invention of random peptides cardiac arrest, edema, pulmonary embolism, glycosylation, iron binding and/or Tf receptor binding. The transferrin fusion proteins are useful for treating, preventing or ameliorating The present invention relates to a library (1) of modified fusion proteins of transferrin (Tf) and therapeutic proteins with increased serum half-life or serum stability. Preferred fusion proteins include those modified so that the Tf moiety exhibits no or reduced pIII protein and the insertion of peptides

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RESULT 5
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Best Local S
Matches 17
The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants of granulocyte colony stimulating factor receptor, glucagon, interlevin 6, platelet-derived endothelial cell growth factor, cyclin-dependent kinase inhibitor 1C, cellular tumour antigen P3, and vasoactive intestinal polypeptide receptor 2. The polypeptides and their associated nucleic acids are useful for identification of variant sequences and detection of candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various disorders including cardiovascular diseases such as arteriosclerosis, myocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as hypertrophy, immune disorders such as immune complex nephritis, multiple sclerosis, cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P53; cyclin-dependent kinase inhibitor 10; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; wyocardial infarction; coronary arterial thrombosis; renal disease; diabetic nephropathy; muscular disease; immune disorder; sarcoidosis; multiple sclerosis; immune complex nephritis; deep vein thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Fig 80; 519pp; English.
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10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Angiotensin converting enzyme (ACEV) splice variant protein #80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200136632-A2
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DB; AAS06080.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      David A,
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99IL-00133455
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Khosravi R,
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1e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bernstein
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JAN-2002; 2002US-0348769P.
18-JAN-2002; 2002US-0350701b.
19-MAR-2002; 2002US-0366659P.
10-MAY-2002; 2002US-0379907P.
                                                                                                                                                                                                                                                                                                           Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                           allergy or anemia), hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                New human metalloproteins and polynucleotides, useful for diagnosing, treating or preventing autoimmune or inflammatory disorders (e.g. AIDS, allergy or anemia), multiple sclerosis, osteoarthritis, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-598523/56.
N-PSDB; ACF35812.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human metalloprotein (MEPR) polypeptide (Id-7509328CD1).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (INCY-) INCYTE GENOMICS
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M, Chien D,
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Yue H,
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The invention relates to novel human metalloproteins (MEPR) and encoding polynucleotides. The human MEPR polypeptides, polynucleotides and modulators are useful for diagnosing, treating or preventing disorders associated with aberrant expression of MEPR, particularly cell proliferative disorders (e.g. arteriosclerosis, atherosclerosis, cirrhosis, hepatitis, paroxysmal nocturnal hemoglobinuria, polycythemia vera, psoriasis, primary thromobocytopenia or cancer), developmental disorders (e.g. renal tubular acidosis, anemia or mental retardation), neurological disorders (e.g. Alzheimer's disease, Parkinson's disease or epilepsy), autoimmune/inflammatory disorders (e.g. AIDS, allergies, asthma, autoimmune thyroiditis, contact dermatitis, Crohn's disease,

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RESULT 7
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Best Local (
                                                                                            The invention relates to novel human metal binding proteins (MBP) and encoding polynucleotides. The MBPs can be expressed by standard recombinant methodology. The polypeptides and polynucleotides are useful in diagnosing, treating and preventing diseases or conditions associated with the decreased expression or overexpression of MBP, such as cell proliferative (e.g. cancer, atherosclerosis), neurological (e.g. AIDS, allergies) and developmental (e.g. Hypothyroidism, Cushing's syndrome) disorders, or infections. These are also useful in assessing the effects of exogenous compounds on the expression of nucleic acid and amino acid sequences of MBP or its fragments are useful in screening compounds for effectiveness as agonist or antagonist of the polypeptides, or in altering the expression of the target polynucleotide and compounds for the captession of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the target polynucleotide and compounds or in altering the expression of the captes of the polypeptides.
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                                 that specifically bind to or modulate the activity of the polypeptide. A microarray comprising different nucleotide molecules affixed at distinct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and preventing expression e.g.
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anticonvulsant; nootropic; r
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The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders, endocrine autoimmune/inflammatory disorders, developmental disorders, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               scnmadt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthsborne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirtol Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, G; Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
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12-SEP-2002;
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Pred. No. 1.8e-05;
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Best Local S
Matches 17
The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliterative disorders, endocrine autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the
                                                                                                                                                                                                                                                                                            New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
                                                                                                                                                                                                                                                                                                                                                                                                           Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder L;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gersti:
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirt,
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, 19
Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                              Claim
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12-SEP-2002;
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DB; ACN42435.
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                                                                                                                                                                                                                                            27; Page; 190pp;
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The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants of granulocyte colony stimulating factor receptor, glucagon, interleukin 6, platelet-derived endothelial cell growth factor, cyclin-dependent kinase inhibitor 1C, cellular tumour antigen P53, and vasoactive intestinal polypeptide receptor 2. The polypeptides and their associated nucleic acids are useful for identification of variant sequences and detection of candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various disorders including cardiovascular diseases such as arteriosclerosis, myocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as hypertrophy, immune disorders such as immune complex nephritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Angiotensin converting enzyme splice variant; ACEV; interleukin 6; granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P53; cyclin-dependent kinase inhibitor IC; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; myocardial infarction; coronary arterial thrombosis; renal disease; diabetic nephropathy; muscular disease; immune disorder; sarcoidosis; multiple sclerosis; immune complex nephritis; deep vein thrombosis; nonarcoidotic pulmonary granulomatous disease; endothelial abnormality; vascular disorder; asbestosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Fig 38; 519pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel alternative aplicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibodies.
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RESULT 11
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P5; cyclin-dependent kinase inhibitor 10; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; myocardial infarction; coronary arterial thrombosis; renal disease; diabetic nephropathy; muscular disease; immune disorder; sarcoidosis; multiple sclerosis; immune complex nephritis; deep vein thrombosis; nonarcoidotic pulmonary granulomatous disease; endothelial abnormality;
                       The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants of granulocyte colony stimulating factor receptor, glucagon, interleukin 6, platelet-derived endothelial cell growth factor, cyclin-dependent kinase inhibitor 1C, cellular tumour antigen P53, and vasoactive intestinal polypeptide receptor 2. The polypeptides and their associated nucleic acids are useful for identification of variant sequences and detection of candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various disorders including cardiovascular diseases such as arteriosclerosis, myocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as the property of the inventor of the inventor of the coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as the property of the coronary arterial thrombosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases such as asbestosis and vascular pathologies involving an endothelial abnormality such as deep vein thrombosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-NOV-1999;
10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Fig 37; 519pp; English.
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DB; AAS06037.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   David A,
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   immune complex nephritis,
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Pred. No.
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es 0;
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multiple sclerosis,
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RESULT 12
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cytostatic; immunosuppressive; antiatherosclerotic; cardiant;
gynaecological; immunostimulant; antianaemic; haemostatic;
antiinflammatory; dermatological; antibacterial; virucide; an
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antiparkinsonian; nootropic; anti-HIV; antiasthmatic; antiallergic;
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Pred. No.
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                                                                                                                                                                                                                                                      cc minus the signal peptide. The invention relates to modified If fusion comprising at least one therapeutic procesin, polypeptide or cc peptide, in which the If portion is engineered to extend the serum half-cc life or bioavailability of the molecule. The modified If fusion protein preferably comprises a human If molecule. The modified If fusion protein cc preferably comprises a human If molecule. The modified to reduce cor prevent glycosylation, iron binding and/or transferrin receptor cc binding, having at least one amino acid substitution, deletion or cc addition in the hinge region, or at residues Asp-63, Gly-65, Tyr-95, Tyr-618, Lys-206, His-207, His-249, Asp-392, Tyr-426, Tyr-514, Tyr-517, His-62, Tyr-95, Tyr-120, Arg-124, Ala-126, Gly-127, Thr-452, Arg-456, Ala-458 and cc Gly-459, or a mutation which prevents glycosylation at Asn-413 or Asn-611 cc (all claimed). Nucleic acids encoding such fusion proteins, vectors, host cc reating a disease or disease symptom, or for delivering a therapeutic cc agent complexed to the ferric iron of transferrin to the inside of a cell cor across the blood-brain barrier. The modified fusion protein, or a cucleic acid encoding it, can be used in the diagnosis, prognosis, cc endocrine, nervous, immune, restriatory, cardiovasqular, repredictive and cc endocrine, nervous, immune, restriatory, cardiovasqular, repredictive and cc endocrine.
                                                                          Query Match
Best Local S
Matches 17
                                                                                                                                                                                     endocrine, nervous, immune, respiratory, cardiovascular, reproductive and digestive systems, diseases and/or disorders relating to the blood or to cell proliferation, inflammatory conditions, and to treat viral, fungal, bacterial or parasitic infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New fusion protein, useful in the diagnosis and treatment of diseases or disorders relating to the respiratory, cardiovascular and digestive systems, comprises a transferrin protein fused to a therapeutic protein.
                                                                                                                                                   Sequence 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 280-281; 298pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-AUG-2001;
30-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is the protein sequence of human transferrin (Tf),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOR-) BIOREXIS PHARM CORP
 251
                                                                          17;
                                     1 VVARSMGGKEDLIWELL 17
                                                                                              Similarity
VVARSMGGKEDLIWELL 267
                                                                                                                                                     Α
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2001US-0334059P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "Iron binding site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Iron binding site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Hinge region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Hinge region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Carbonate ion binding
                                                                          0
                                                                                            Score 86;
Pred. No.
                                                                          Mismatches
                                                                   DB 6; bc. 2e-05; 0;
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                                                                                                            Length 679;
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RESULT 13

RESULT 14 ADK15869

RXU

ADK15869;

ADK15869 standard; protein; 679

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                                                                                                                                                                            This invention relates to a novel fusion protein which comprises a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region. The invention may be useful for the development of compounds with cytostatic, antibacterial, virucide, antiparasitic, immunosuppressive or antiarthritic activity. In addition, the sequences disclosed may be useful for gene therapy. The fusion protein is useful for preparing a composition for treating a disease or disease symptom in a patient for example septic shock, endotoxic shock, cachexia syndromes associated with bacterial, viral or parasitic infections, neoplasm, autoimmune disease, arthritis or adverse effects associated with treatment for preventing graft rejection. The present sequence is that of the mature human transferrin protein which is related
                                                                                                                                                                                                                                                                                                                                                                                                                                   New fusion protein comprising a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region, useful for preparing a composition for treating e.g., septic shock, neoplasm or autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibody variable region; cytostatic; antibacterial; virucide;
antiparasitic; immunosuppressive; antiarthritic; gene therapy;
septic shock; endotoxic shock; cachexia syndrome; bacterial infection;
                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; ADH89358.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-022093/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sadeghi H,
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30-NOV-2001; 2001US-0334059.
30-AUG-2002; 2002US-00231494.
30-AUG-2002; 2002US-0406977P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     viral infection; parasitic infection; neoplasm; autoimmune disease; arthritis; graft rejection; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human transferrin protein mature amino acid sequence
                                                                                                                                                                 to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 3; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fusion protein; transferrin protein; glycosylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH89360 standard; protein;
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                 1 VVARSMGGKEDLIWELL 17
VVARSMGGKEDLIWELL 267
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                                                              Conservative
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                                                                              100.0%; Score 86;
100.0%; Pred. No.
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                                                                                 2e-05;
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                                                                                               Length
                                                                  Indels
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RESULT 15
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XX ADL70
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XX Inmunr
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                                                                                              Immunosuppressive; Haemostatic; Antiallergic; Antiasthmatic; Dermatological; Antiinflammatory, Antibacterial; Vasotropic; Neuroprotective; Vulnerary; Nephrotropic; Neuroprotective; Cytostatic; Cerebroprotective; Vulnerary; Antiparkinsonian; Nootropic; Cardiant; Antianaemic; Antiatherosclerotic; Thrombolytic; Anticoagulant; Gastrointestinal; Respiratory; transferrin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises a fusion protein that contains a modified transferrin (Tf) protein fused to a therapeutic protein/peptide (e.g. beta-interferon - IFN, glucagons-like peptide - GLP-1, erythropoietin mimetic peptide - EMP1, T-20 and soluble toxin receptor). The fusion protein of the invention is useful for epitope tagging. The present amino acid sequence represents the mature human Tf protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-AUG-2001; 2001US-0315745P.
30-NOV-2001; 2001US-0334059P.
30-AUG-2002; 2002US-00231494.
30-AUG-2002; 2002US-0406977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fusion protein; transferrin; Tf; beta-interferon; beta-IFN; glucagons-like peptide; GLP-1; erythropoietin mimetic peptide; T-20; soluble toxin receptor; epitope tagging; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 679
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 11-MAR-2004
                                                                                                                                                                                  Human transferrin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prior
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fusion protein comprising a modified transferrin (Tf) protein fused therapeutic protein or peptide, useful for epitope tagging.
                                                       sapiens
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 3; 70pp; English.
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                                                                                                                                                                                  Tf, mature protein SEQ ID
                                                                                                                                                                                                                                                                   protein; 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sadeghi H,
                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                  protein;
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Pred. No.
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2e-05;
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                                                                                                                                                                          CC blood. The modified fusion proteins are useful in diagnosis, prognosis, considered fusion proteins are useful in diagnosis, prognosis, considered fusion and/or treatment of autoimmune disorders; diseases and cc disorders of haematopoietic cells (e.g., leukopenia, neutropenia, anaemia cc and thrombocytopenia); allergic reactions such as allergic asthma, cc anaphylaxis, IgE-mediated allergic reactions such as allergic asthma, rhinitis and cerema; inflammatory conditions e.g., inflammation associated with cc infection (e.g., septic shock, sepsis), ischamenia-reperfusion injury, cc mephritis, Crohn's disease, multiple sclerosis, respiratory disorders (asthma and allergy), gastrointestinal disorders (inflammatory bowel conditions of e.g., gastric, ovarian, lung, bladder), CNS disorders (multiple sclerosis, stroke, traumatic brain injury, neurodegenerative disorders such as Parkinson's disease, Alzheimer's disease), etc. The cusion protein is also useful as an adjuvant to enhance antibacterial or canifungal immune responses, antiparasitic immune responses, etc. The cundetermined significance (MGUS), Waldenstrom's disease, plasmacytomas, cc adult respiratory distress syndrome, for stimulating wound repair, for condetermined algorithmis infections of joints, bones, skin, etc. The fusion protein is also useful for treating or preventing thrombosis, myocardial collemention, cancers, thrombocytopenia, sickle cell anaemia,
                                                                                    Matches
                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a library (I) of modified fusion proteins of transferrin (Tf) and therapeutic proteins with increased serum half-life or serum stability. Preferred fusion proteins include those modified so that the Tf molety exhibits no or reduced glycosylation, iron binding and/or Tf receptor binding. The transferrin fusion proteins are useful for treating, preventing or ameliorating disorders or diseases of endocrine system, nervous system, immune system, respiratory system, cardiovascular system, diseases and/or disorders relating to cell proliferation, and/or diseases or disorders relating to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel library containing several fusion proteins each of which comprises first transferrin polypeptide fused to at least one second peptide, useful for screening for transferrin fusion protein having the particular
                                                                                                                                                                                                                 glomerulonephritis, cardiac arrest, edema, pulmonary embolism, atherosclerosis, etc. The present sequence is human Tf mature used to make the modified Tf fusion proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-AUG-2002; 2002US-0406977P
10-MAR-2003; 2003US-00384060
09-JUL-2003; 2003US-0485404P
                                                                                                                                                                          Sequence
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86
 BLOSUM62
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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                                                                                                                                                                                                                           transferrin - pig
lactoferrin precur
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transferrin - mous
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hypothetical prote
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transferrin precur
transferrin precur
lactotransferrin p
                  quinone oxidoreduc ABC-type transport ABC transporter, A
                                                                   receptor protein-t
embryo kinase 5 -
hypothetical ORF-1
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 hypothetical prote
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ALIGNMENTS

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transferrin precursor - rabbit

C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 30-Sep-193 #sequence revtsion 30-Sep-193 #text_change 22-Jun-1999
C;Accession: S16246; A61239; C61573; S00335; S02694; A26504; S14853
R;Banfiteld, D.K.; Chow, B.K.C.; Funk, W.D.; Robertson, K.A.; Umelas, T.M.; V
Biochim. Biophys. Acta 1089, 262-265, 199; Robertson, K.A.; Umelas, T.M.; V
A;Title: The nucleotide sequence of rabbit liver transferrin cDNA.
A;Reference number: S16246; MUID:91274362; PMID:2054387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Map position: 3q21-3q21
A;Introns: 15/1; 72/3; 119/1; 168/1; 212/2; 231/1; 290/3; 350/1; 401/3; 433/1; 444/1; 49
C;Function:
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A; Residues: 564-698 < RE4 >
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A;Accession: I68160
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A; Residues: 1-72 < RE3 >
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A;Residues: 636-696 KRZ>
A;Residues: 636-696 KRZ>
A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:Gene 56, 109-116, 1987
Gene 56, 109-116, 1987
A;Title: Complete structure of the human transferringene. Comparison with analogous chi
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Proc. Natl. Acad. Sci. U.S.A. 86, 7260-7264, 1989
A;Title: Changes in brain gene expression shared by scrapie and Alzheimer disease.
A;Reference number: 148174; MUID:89386721; PMID:2780570
A;Accession: 16133
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A;Title: A cloned gene for human transferrin.
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A;Title: The complete amino acid sequence of human serum transferrin.
A;Reference number: A93911; MUID:8222166; PMID:6953407
A;Contents: annotation; disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GDB:120432; OMIM:190000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;356-686/Domain: transferrin repeat homology <TRH2>
;28-67,38-58,137-213,156-350,177-193,180-196,190-198,246-260,358-615,364-396,374-387,42
;432,630/Binding site: carbohydrate (Asn) (covalent) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;20-698/Product: transferrin #status experimental <MAT>;20-350/Domain: transferrin repeat homology <TRH1>
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A; Cross-references: UNIPARC: UPI00001744AA
C; Superfamily: transferrin; transferrin repeat homology
C; Superfamily: transferrin; transferrin repeat homology
C; Keywords: duplication; glycoprotein; iron transport; metal binding; plasma
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-694/Product: transferrin #status experimental <MAT>
F;19-349/Domain: transferrin repeat homology <TRH1>
F;35-568/Domain: transferrin repeat homology <TRH2>
F;35-682/Domain: transferrin repeat homology <TRH2>
F;37-66,37-7,136-212,155-349,176-192,179-195,189-197,245-259,357-611,363-395,373-386,420
F;508/Binding site: carbohydrate (Asn) (covalent) #status experimental
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A; Residues: 482-515,'V',517-544 <EVA>
A; Residues: 482-515,'V',517-544 <EVA>
A; Rote: 516-IIe was also found
A; Note: 516-IIe was also found
R; Heaphy, S.; Williams, J.
Biochem. J. 205, 611-617, 1982
A; Title: The preparation and partial characterization of
A; Reference number: A26504; MUID:83074540; PMID:6816218
R;Roush, E.D.; Fierke, C.A.
Biochemistry 31, 12536-12542, 1992
A;Title: Purification and characterization
A;Reference number: I47228; MUID:93099129;
                                                                                                                                  carbonic anhydrase II inhibitor (transferrin homolog) precursor - pig C;Species: Sus scrofa domestica (domestic pig) C;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 09-Jul-2004 C;Accession: 147228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Accession: A26504
A;Molecule type: protein
A;Residues: 19-24,'N',26,'X',28-29,'S' <HEA>
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A;Title: Evidence for a single glycan moiety in rabbit serum transferrin A;Reference number: S02694; MUID:89005676; PMID:3169252
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Biol. Chem. Hoppe-Seyler 369, 93-96, 1988
Biol. Chem. Hoppe-Seyler 369, 93-96, 1988
A;Title: Isolation, characterization and N-terminal amino-acid sequence of rabbit transfe A;Reference number: S00335; MUID:88209278; PMID:3365331
A;Accession: S00335
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A; Residues: 19-26, 'X', 28-36, 'X', 38-53 < CHU>
A; Cross-references: UNIPARC: UPI00001744A7
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R;Chung, M.C.M.; Chan, S.L.; Shimizu, S.
Int. J. Biochem. 23, 609-616, 1991
A;Title: Purification of transferrins and lactoferrin using A;Reference number: A61573; MUID:91293379; PMID:2065820
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Cell. Immunol. 134, 225-234, 1991
A;Title: Iron carrier proteins fa
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A; Cross-references: UNIPARC: UPI00001744A8
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A: Residues: 19-36 < PIE>
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Pred. No. 0.00012;
1; Mismatches 1
             of a carbonic anhydrase PMID:1463741
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A;Residues: 1-144,'V',146-163,'PP',166-339,'A',341-438,'Y',440-513,'R',515-708 < A;Cross-references: UNIPARC:UPI000016C335; EMBL:X57084; NID:g505; PIDN:CAA40366. R;Pierce, A.; Colavizza, D.; Benaissa, M.; Maes, P.; Tartar, A.; Montreuil, J.; Eur. J. Biochem. 196, 177-184, 1991. A;Title: Molecular cloning and sequence analysis of bovine lactotransferrin. A;Reference number: S14110; MUID:91160550; PMID:2001696
                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-65, 'PG', 'B-296, 'S', 298-339, 'A', 341-708 <GOO>
A; Cross-references: UNIPARC: UPI0000174495; GB: M63502
A; Note: the authors translated the codon CCG for residue 6
R; Mead, P.E.; Tweedie, J.W.
Nucleic Acids Res. 18, 7167, 1990
A; Title: CDNA and protein sequence of bovine lactoferrin.
A; Reference number: S13097; MUID: 91088328; PMID: 2263492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Goodman, R.E.; Schanbacher, F.L.
Biochem. Biophys. Res. Commun. 180, 75-84, 1991
Biothem. Bovine Lactoferrin mRNA: sequence, analysis, and
A;Reference number: JT0595; MUID:92028986; PMID:1718281
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                                 A;Molecule type: mRNA
A;Residues: 28-33,'DS',36-38,'P',40-708 <MF
A;Cross-references: UNIPARC:UPI0000174496;
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A;Residues: 20-35;82-114;148-163,'PP',166-178,'V','P';183-190;205-212;230-239;304-339;5:
A;Cross-references: UNIPARC:UPI000017448C; UNIPARC:UPI000017448D; UNIPARC:UPI000017448E
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A;Residues: 3-144,'V',146-339,'A',341-438,'Y',440-513,'R',515-708 <PI2>
A;Cross-references: UNIPARC:UPI000017448B; EMBL:X57084
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A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-708 <TSA>
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C;Date: 31-Mar-1992 #sequence revision 21-Nov-1997 #text_change 09-Jul-2004
C;Accession: 145919; S14674; S14110; S18517; JT0595; S13097; S18518; S13881;
R;Tsang, T.C.; Burns, D.K.; Wang, F.; Pan, Y.
FASEB J. 6, 233, 1991
A;Title: Cloning of a 80-kD advanced glycosylation end product (AGE) binding
A;Reference number: 145919
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A;Cross-references: UNIPARC:UPI0000177C50
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A;Molecule type: protein
A;Residues: 20-25;302-308;359-366,'X',368-376,'X',378 <SHI>A;Residues: 20-25;302-308;359-366,'X',368-376,'X',378 <SHI>A;Cross-references: UNIPARC:UPI00001744A1; UNIPARC:UPI00001744A2; UNIPARC:UPI00001744A3
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron; iron binding; metalloprotein; milk
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-708/Product: lactotransferrin #status experimental <MAT>
F;20-355/Domain: transferrin repeat homology <TRH1>
                                                                                                C;Accession: JC2323
R;Le Provost, P.; Nocart, M.; Guerin, G.; Martin, P. Biochem. Biophys. Res. Commun. 203, 1324-1332, 1994
A;Title: Characterization of the goat lactoferrin cDNA:
A;Reference number: JC2323; MUID:94380047; PMID:8093048
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A;Cross-references: UNIPARC:UPI00000475A8
R;Shimazaki, K.; Tanaka, T.; Kon, H.; Oota, K.; Kawaguchi,
J. Dairy Sci. 76, 946-955, 1993
A;Title: Separation and characterization of the C-terminal
A;Reference number: A56659, MUID:93253156; PMID:8486845
A;Accession: A56659
                                                                                                                                                                                                                                                                                  C;Species: Capra aegagrus hircus (domestic goat)
C;Date: 20-Feb-1995  #sequence_revision 20-Feb-1995  #text_change 07-May-1999
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A;Title: Identification of the bactericidal domain of lactoferrin.
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A; Residues: 20-27, 'X', 29-37, 'X', 39-54, 'X', 56-59 <REJ>
A; Cross-references: UNIPARC: UPI00001744A0
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Comp. Biochem. Physiol. B 93, 929-934, 1899
A;Title: Purification and characterization of bovine lactoferrin
A;Reference number: PL0148; MUID:90031466; PMID:2805645
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A;Residues: 28-38,'P',40-86,'C',88-708 <ME3>
A;Cross-references: UNIPARC:UPI000017449F; EMBL:X54801
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A;Cross-references: UNIPARC:UPI0000174493; UNIPARC:UPI0000174497; UNIPARC:UPI0000174498;
A; Molecule type: mRNA
A; Residues: 1-708 < LEP >
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    F;36-60/Region: antimicrobial F;359-696/Domain: transferrin repeat homology <TRH2>
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                                                                     A; Accession: JC2323
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                                                                                                                                                                                                                                                                                                                                                                    lactoferrin - goat
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Pred. No.
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C;Species: Equus (aballus (domestic horse)
C;Species: Equus (aballus (domestic horse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: S33761; S02145;
R;Carpenter, M.A.; Broad, T.E.
Biochim. Biophys. Acta 1173, 230-232, 1993
A;Title: The cDNA sequence of horse transferrin.
A;Reference number: S33761; MUID:93277958; PMID:8504171
A;Reference number: S33761
A;Accession: S03176
A;Accession: S03176
A;Accession: S03176
A;Accession: S03176
A;Title: A growth-promoting factor for human myeloid leukemia cells from horse serum ide
A;Reference number: S02145; MUID:89076897; PMID:2909248
A;Accession: S03176
A;Molecule type: mRNA
A;Residues: 1-695 <ESC>
A;Cross-references: UNIPROT:P12346; UNIPROT:Q63602; UNIPARC:UPI0000167A8E; EMBL:X77158;
R;ESCTIVA, H.; Pierce, A.; Coddeville, B.; Gonzalez, F.; Benaissa, M.; Leger, D.; Wierus
Biochem. J. 307, 47-55, 1995
                                                                                                                                                                                                                                                                                                        C:Species: Rattus norvegicus (Norway rat)
C:Date: 16-Fab-1995 #sequence voision 12-May-1995 #text change 09-Jul-2004
C:Date: 16-Fab-1995 #sequence revision 12-May-1995 #text change 09-Jul-2004
C:Accession: S49163; S54980; A30014; A14679; A53289; A30512; I52203
R:Escriva, H.; Pierce, A.; Coddeville, B.; Gonzalez, F.; Benaissa, M.; Leger, submitted to the EMBL Data Library, January 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Accession: $02145
A;Molecule type: protein
A;Residues: 20-35,'X',37,'X',39-40,'X',43-44 <YOS>
A;Residues: 20-35,'X',37,'X',39-40,'X',43-44 <YOS>
A;Cross-references: UNIPARC:UPI0000177C56
C;Complex: monomer
C;Superfamily: transferrin transferrin repeat hom
C;Keywords: duplication; glycoprotein; iron transp
F;1-19/Domain: signal sequence #status predicted <
F;20-706/Product: transferrin #status experimental
                                                                                                                                                                                                                                               A; Description: Rat mammary gland A; Reference number: S49163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transferrin precursor - rat N; Alternate names: lung-derived growth factor; siderophilin
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                                                                                                                                                                                                           A; Accession: S49163
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;1-19/Domain: signal sequence #status predicted <SIG>
;20-706/Product: transferrin #status experimental <AMT>
;358-694/Domain: transferrin #status experimental <AMT>
;358-694/Domain: transferrin repeat homology <TRH2>
;26-64,36-55,134-215,174-190,177-198,187-200,248-262,360-623,366-398,376-389,423-701,44
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                                                                                                                                                                       Status: preliminary
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                                                                                                                                                                                                                                                                           transferrin: glycan structure,
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Pred. No. 0.002;
1; Mismatches
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Pred. No. 0.
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A;Cross-references: UNIPARC:UPI0000177C51
R;Cavanaugh, P.G.; Nicolson, G.L.
J. Cell. Biochem. 47, 261-271, 1991
A;Title: Lung-derived growth factor that stimulates the growth A;Reference number: A53289; MUID:92165927; PMID:1791188
A;Accession: A53289
                                                                                                                                                                                                                                                                                  C;Superfamily: transferrin; transferrin repeat homology C;Keywords: duplication F;20-348/Domain: transferrin repeat homology <TRH1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Aldred, A.R.; Howlett, G.J.; Schreiber, G.
Biochem. Biophys. Res. Commun. 122, 960-965, 1984
A;Title: Synthesis of rat transferrin in Escherichia coli containing a recombinant bacter
A;Reference number: I52203; MUID:84307580; PMID:6236811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 20-30;639-643,'KD',646,'LKACD' <PUR>
A;Cross-references: UNIPARC:UPI0000177C52; UNIPARC:UPI0000177C53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Note: sequence modified after extraction from NCBI backbone
A;Note: sequence extracted from NCBI backbone (NCBIP:86115, NCBIP:86116, NCBIP:86114)
R;Purves, L.R.; Purves, M.; Linton, N.; Brandt, W.; Johnson, G.; Jacobs, P.
Biochim. Biophys. Acta 966, 318-327, 1988
A;Title: Properties of the transferrin associated with rat intestinal mucosa.
A;Reference number: A30512; MUID:88327006; PMID:3046665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: protein
A;Residues: 89,'Y',91'A',93-99,'V',101-102;'N',233,'AN',236-243;401-406,'N',408 <CAV>
A;Cross-references: UNIPARC:UPI00000E6F28; UNIPARC:UPI00000E76D0; UNIPARC:UPI00000E7E7F
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A;Title: The synthesis and secretion of rat transferrin.
A;Reference number: A14679; MUID:80049855; PMID:500689
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A;Tille: Transferrin messenger ribonucleic acid: molecular cloning and hormonal regulatic A;Reference number: A30014; MUID:87053639; PMID:3023031
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A;Residues: 7-25,'X',27-56,'A',58-64,267-295 <RES>
A;Cross-references: UNIPARC:UPI00000E5C48; GB:M26113; NID:g207437; PIDN:AAA42266.1; PID:E
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A;Reference number: S54980; MUID:95234054; PMID:7717992
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                                                                                                                                             Matches
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                   Local
   270
                                                                    1 VVARSMGGKEDLIWELL 17
                                                                                                                                                                                   Similarity
                                                                                                                                             Conservative
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76.5%;
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Pred. No.
                                                                                                                                                    Mismatches
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N;Alternate names: lactoferrin
C;Species: Homo sapiens (man)
C;Date: 31-Mar-1992 #sequence_

#sequence_revision

21-Nov-1997 #text_change 08-Dec-2000

lactotransferrin precursor [validated]

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A;Accession: $20841

A;Molecule type: protein

A;Residues: 20-28; X',30-31 <$T2>
A;Residues: 20-28; X',30-31 <$T2>
A;Cross-references: UNIPARC:UDI0000174487

R;Rado, T.A.; Wei, X.; Benz Jr., E.J.

Blood 70, 989-993, 1987

A;Title: Isolation of lactoferrin cDNA from a human myeloid library and expression of mR
A;Title: Isolation of lactoferrin cDNA from a human myeloid library and expression of mR
A;Reference number: $07160; MUID:88001031; PMID:3477300

A;Accession: $07160

A;Molecule type: mRNA
A;Residues: 436-487,'A',489-711 <RADD
A;Cross-references: UNIPARC:UPI000016ABE2; EMBL:M18642; NID:g186815; PIDN:AAA86665.1; PI
R;Panella, T.J.; Liu, Y.; Huang, A.T.; Teng, C.T.
Cancer Res. 51, 3037-3043, 1991

A;Title: Polymorphism and altered methylation of the lactoferrin gene in normal leukocyt
A;Beference number: A61169; MUID:91235214; PMID:1674448

A;Cross-references: A61169; MUID:91235214; PMID:1674448
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A;ACCEBBLUM.
A;Molecule type: mRNA
A;Residues: 3-711 <POW>
A;Residues: 3-711 <POW>
A;Residues: 3-711 <POW>
A;Cross-references: UNIPARC:UDIO0016AC28; EMBL:X52941; NID:g34411; PIDN:CAA37116.1;
A;Cross-references: UNIPARC:UDIO0016AC28; EMBL:X52941; NID:g34411; PIDN:CAA37116.1;
B:Stowell, K.M.; Rado, T.A.; Funk, W.D.; Tweedie, J.W.
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A; Molecule type: mRNA
A; Residues: 1-148, 'T', 150-422, 'C', 424-711 <REY>
A; Residues: 1-148, 'T', 150-422, 'C', 424-711 <REY>
A; Cross-references: UNIPARC: UPI000015C582; EMBL: X53961; NID: g34415; PIDN: CAA37914.1;
A; Teng, C.T.; Liu, Y.; Yang, N.; Walmer, D.; Panella, T.

Wol. Endocrinol. 6, 1969-1981, 1992
A; Title: Differential molecular mechanism of the estrogen action that regulates lactor A; Reference number: A45401; MUID: 93125571; PMID: 1480183
A; Accession: A45401
                                                                                                  R;Stowell, K.M.; Rado, T.A.; Funk, W.D.; Tweedie, J.W. Biochem. J. 276, 349-355, 1991
A;Title: Expression of cloned human lactoferrin in baby-hamster kidney A;Reference number: S15853; MUID:91264786; PMID:2049066
A;Accession: S15853
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A,Note: sequence extracted from NCBI backbone (NCBIP:122202)
R,Powell, M.J.; Ogden, J.E.
Nucleic Acids Res. 18, 4013, 1990
A,Title: Nucleotide sequence of human lactoferrin cDNA.
A,Reference number: $10324, MUID:90326549, PMID:2374734
A,Recession: $10324
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Nucleic Acids Res. 18, 5288, 1990.
A;Title: Complete nucleotide sequence of human mammary gland lactoferrin.
A;Reference number: S11228; MUID:90384839; PMID:2402455
                                                                                                                                                                                    A,Experimental source: normal breast tissue R,Metz-Boutigue, M.H.; Jolles, J.; Mazurier, Eur. J. Biochem. 145, 659-666, 1984
                                                                      A; Reference number: A31000;
A; Accession: A31000
                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 3-701, 'SWKPVN' < PAN>
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A; Residues: 20-31 <ST1>
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A; Residues: 1-15 < TEN>
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20-140, 142-169, 171-203, 'L', 205, 'K', 207-208, 'K', 210-385, 'Q', 387-391, 'W', 393
                                           protein
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PMID:6510420
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                                                                                                                                                                                                                           J.; Schoentgen,
                                                                                                                                                     structural comparisons with
                                                                                                                                                                                                                       F.; Legrand, D.;
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                                                                                                                                                                                                                               Spik,
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A;Molecule type: protein
A;Residues: 1-8, 'X',10-18, 'XE' <CHU>
A;Cross-references: UNLTBARC:UPI0000174486
A;Cross-references: UNLTBARC:UPI0000174486
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron transport; |
F;1-659/Product: transferrin repeat homology <TRH1>
F;1-335/Domain: transferrin repeat homology <TRH1>
                                                                                                                                                                                                                                                  A;Note: the authors suggest transferrin from gastric mucosa R;Chung, M.C.M.; Chan, S.L.; Shimizu, S. Int. J. Biochem. 23, 609-616, 1991
A;Title: Purification of transferrins and lactoferrin using A;Reference number: A61573; MUID:91293379; PMID:2065820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:P09571; UNIPARC:UPI00001130C2; EMBL:X12386; NID:g2126; PIDA A;Note: 308-Arg was also found R;Baldwin, G.S.; Bacic, T.; Chandler, R.; Grego, B.; Pedersen, J.; Simpson, R.J.; Toh, Comp. Biochem. Physiol. B 95, 261-268, 1990 A;Title: Isolation of transferrin from porcine gastric mucosa: comparison with porcine A;Reference number: A60520; MUID:90227903; PMID:2328566
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C;Superfamily: transferrin; glycoprotein; iron binding; milk
C;Keywords: duplication; glycoprotein; iron binding; milk
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-711/Product: lactotransferrin #status experimental <MAT>
F;21-356/Domain: transferrin repeat homology <TRH1>
F;360-699/Domain: transferrin repeat homology <TRH2>
F;360-699/Domain: transferrin repeat homology <TRH2>
F;360-699/Domain: transferrin repeat homology <TRH2>
F;360-897/S91-801/S1-218,177-193,190-201,251-265,503-697,595-609/Disulfide bonds: #status
F;358-400,378-391,425-706,447-669,479-554,513-527,524-537,647-652/Disulfide bonds: #sta
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A;Note: this is the final paper in a series
R;Houen, G.; Hoegdall, E.V.; Barkholt, V.; Norskov, L.
Eur. J. Biochem. 241, 303-308, 1996
A;Title: Lactoferrin: similarity to diamine oxidase and A;Reference number: S74119; MUID:97054624; PMID:8898921
A;Accession: S74119
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-8,'X',10-11,'X',13-15 <BA2>
A;Cross-references: UNIPARC:UPI0000174485
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Nucleic Acide Res. 16, 8720, 1988
A;Title: Nucleotide sequence of porcine liver transferrin.
A;Reference number: S01384; MUID:88335629; PMID:3419934
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A;Residues: 'G',23-24,'R',26-27,'XX',30-32 <HOU>
A;Cross-references: UNIPARC:UPI000017448A
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C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999
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Pred. No. 0.0044;
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C.Species: Mus musculus (house mouse)
C.Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C.Accession: A28438; A41205
R.Pentecost, B.T.; Teng, C.T.
J. Biol. Chem. 262, 10134-10139, 1987
A.Title: Lactotransferrin is the major estrogen inducible protein of mouse in A.Reference number: A92596; MUID:87280033; PMID:3611056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F;36-48/Region: antimicrobial
F;354-691/Domain: transferrin repeat homology <TRH2>
F;254-691/Domain: transferrin repeat homology <TRH2>
F;28-62,38-63,129-212,171-187,184-195,245-259,362-394,372-385,419-698,439-661,471-546,49
F;77,107,206,267/Binding site: iron (Asp. Tyr. Tyr. His) #status predicted
F;135/Binding site: carbonate (Arg) #status predicted
F;135/Binding site: carbonate (Arg) #status predicted
F;497/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;490/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                     lactoferrin precursor - mouse N;Alternate names: lactotransferrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-11, 'W',13-50,'I',52-84,'G',86-120,'L',121-130,'I',132-282,'S',284-571,'Q'
A; Cross-references: UNIPARC:UPI000016C6E2; EMBL:M92089; NID:g164613; PIDN:AAA31102.1; P
A; Experimental source: mammary gland
C; Superfamily: transferrin; transferrin repeat homology
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Biochim. Biophys. Acta 1132, 97-99, 1992
A;Title: Nucleotide and primary amino acid sequence of porcine lactoferrin. A;Reference number: S24173; MUID:92379101; pMID:1511016
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C;Date: 17-Feb-1994 #sequence_revision 17-Feb-1994 #text_change 09-Jul-2004
C;Accession: A45543; S24173
R;Alexander, L.J.; Levine, W.B.; Teng, C.T.; Beattie, C.W.
Anim. Genet. 23, 251-256, 1992
A;Title: Cloning and sequencing of the porcine lactoferrin cDNA.
A;Reference number: A45543; MUID:92367939; PMID:1503259
A;Accession: A45543
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A;Molecule type: mRNA
A;Residues: 1-703 <ALE>
                          A; Molecule type: mRNA
A; Residues: 3-707 < PEN>
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A;Cross-references: UNIPARC:UPI00001744A4; EMBL:J03298
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;Keywords: duplication; glycoprotein; iron; iron binding; metalloprotein; milk
;1-19/Domain: signal sequence #status predicted <SIG>
;20-703/Product: lactoferrin #status predicted <MAT>
;20-30/Domain: transferrin repeat homology <TRH1>
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Best Local S
Matches 13
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Pred. No. 0.
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Pred. No. 0.0064;
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0.0064;
hes 2; Indels
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A;Molecule type: mRNA
A;Roslecule type: mRNA
A;Roslecule type: mSNA
A;Cross-references: UNIPROT:Q92111; UNIPARC:UPI0000E773F; UNIPARC:UPI000016D098; UNIPARC
A;Cross-references: UNIPROT:Q92111; UNIPARC:UPI0000E773F; UNIPARC:UPI000016D098; UNIPARC
A;Note: the authors translated the codon ACC for residue 61 as Tyr, TCC for residue 62 as
R;Idzerda, R.L.; Behringer, R.R.; Theisen, M.; Huggenvik, J.I.; McKnight, G.S.; Brinster,
Mol. Cell. Biol. 9, 5154-5162, 1989
A;Title: Expression from transferrin gene promoter in transgenic mice.
A;Reference number: A33482; MUID:90097932; PMID:2601714
A;Accession: A33482
                                                                                                                                                                                                                                                                                                                                               A;Molecule type: DNA
A;Residues: 1-14;302-311 <IDZ>
A;Residues: 1-14;302-311 <IDZ>
A;Cross-references: UNIPARC:UPI0000177C57; UNIPARC:UPI0000177C58; GB:M30819; GB:M30820
R;Pentecost, B.T.; Teng, C.T.
J. Biol. Chem. 262, 10134-10139, 1987
A;Title: Lactotransferrin is the major estrogen inducible protein of mouse uterine sec: A;Reference number: A92596; MUID:87280033; PMID:3611056
A;Accession: B28438
                                                                                                                                                                                                         A;Molecule type: mRNA
A;Residues: 86-216 <PEN>
A;Cross-references: UNIPARC:UPI000016CEB1; GB:J03299; GB:J02737; NID:g198847; PIDN:AAA394
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication
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C;Date: 30-Jun-1989 #sequence revision 05-Dec-1997 #text_ch
C;Accession: A28446; A33482; B28438
R;Chen, L.H.; B188ell M.J.
J. Biol. Chem. 262, 17247-17250, 1987
J. Biol. Chem. 262, 17247-17250, 1987
A;Title: Transferrin mRNA level in the mouse mammary gland
A;Reference number: A28446; MUID:88086992; PMID:3693348
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F;20-707/Product: lactotransferrin #status predicted <MAT>
F;358-695/Domain: transferrin repeat homology <TRH2>
F;494/Binding site: carbohydrate (Asn) (covalent) #status predicted
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J. Biol. Chem. 266, 21880-21885, 1991
A;Title: Characterization of estrogen-responsive mouse lactoferrin promoter.
A;Reference number: A41205; MUID:92042099; PMID:1939212
A;Accession: A41205
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C;Superfamily: transferrin; transferrin repeat homolo
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A; Residues: 1-15 <LIU>
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Pred. No.
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Pred. No. 0.33;
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and

hypothetical protein PAB1598 - Pyrococcus abyssi (strain Orsay)
C;Species: Pyrococcus abyssi
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C;Accession: A75097

A75097 RESULT 13

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A;Accession: G72464
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-510 <KAW>
A;Cross-references: UNIPF
                                                                                                                   A; Account type: DNA
A; Molecule type: DNA
A; Residues: 1-2958 <BEN>
A; Cross-references: UNIPROT: Q12150;
A; Cross-references: gtrain $288C
                                                                                                                                                                                                                                                                probable membrane protein YLR087c - yeast (Saccharomyces cerevisiae)
N;Alternate names: hypothetical protein L2506
C;Species: Saccharomyces cerevisiae
C;Date: 01-Aug-1995 #sequence_revision 24-May-1996 #text_change_09-Jul-2004
C;Accession: S64921; S64919
R;Benes, V; Rechmann, S; Nentwich, U; Schwager, C; Ansorge, W; Voss, H.
R;Benes, V; Rechmann, S; Nentwich, U; Schwager, C; Ansorge, W; Voss, H.
submitted to the Protein Sequence Database, May 1996
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C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Aeropyrum pernix
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 05-Oct-2004 .
C;Accession: G72464
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A;Tille: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropy A;Reference number: A72450; MUID:99310339; PMID:10382966
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A;Molecule type: DNA
A;Residues: 1-459 <KAW>
A;Cross-references: UNIPROT:Q9UZH9; UNIPARC:UPI00000633CD; GB:AJ248286; GB:AL096836; NID
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A;Accession: A75097
                       A; Reference number: S64899
A; Accession: S64919
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A; Accession: S64921
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                                                                     submitted to the Protein Sequence Database, May 1996
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Molecule type: DNA
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Best Local
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Similarity 52.4%;
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Pred. No. 1
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F;1219-1235/Domain: transmembrane #status predicted F;1928-1944/Domain: transmembrane #status predicted F;2654-2670/Domain: transmembrane #status predicted
                                                                                                                                                                                      C;Superfamily: Saccharomyces cerevisiae probable membrane protein YLR087c C;Keywords: transmembrane protein F;21-37/Domain: transmembrane #status predicted <TM1>
                                                                                                                                                                                                                                                                                             A;Residues: 2353-2958 <POH>
A;Cross-references: UNIPARC:UPI000017904C; EMBL:Z73259; MIPS:YLR087c
A;Experimental source: strain S288C
C;Genetics:
                                                                                                                                                                                                                                          A;Cross-references: SGD:S0004077
A;Map position: 12R
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1653
                                1 VVARSMGGKEDLIWELL 17
LMSRTVGGRVSLLWENL
                                                                   Conservative
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Search completed: June 13, 2006, 04:20:34 Job time : 18.25 secs THIS PAGE BLANK (USPTO)

Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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Result
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1: uniprot_sprot:*
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86
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Q9GL93 INIGE
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ICA PIG
Q7TSX8 MARMO
TREL_CAMDR
Q6LC78_BOVIN
Q6LB7_BOVIN
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07tsx8
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Q9GL95 PHOPH
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RESULT 3
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Pongo pygmaeus (Orangutan).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-FEB-2006, entry version 6. Hypothetical protein DKFZp459H0229. Name=DKFZp459H0229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OSR9L7; PONPY PRELIMINARY; PRT; 697 AA.
OSR9L7;
OSR9L7;
21-DEC-2004, integrated into UniProtKB/TrEMBL
21-DEC-2004, sequence version 1.
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SMR;
GO; G
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MEDLINE=20481910; PubMed=11027333; DOI=10.1073/pnas.97.21.11343;
Cassens I. Vicario S., Waddell V.G., Balchowsky H., Van Belle D.,
Ding W., Fan C., Mohan L., Simoes-Lopes P.C., Bastida R., Meyer A.
Stanhope M.J., Milinkovitch M.C.;
                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                  Bloecker H., Boecher M., Brandt P., Mewes H.W., Weil B., Osanger A., Fobo G., Han M., Wiemann S.; Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001156; Peptidase S60. PANTHER; PTHR11485; Peptidase S60; 1. Pfam; PF00405; Transferrin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stanhope M.J., Milinkovitch M.C.; "Independent adaptation to riverine habitats allowed survival of ancient cetacean lineages."; Proc. Natl. Acad. Sci. U.S.A. 97:11343-11347(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phocoenoides phocoena (Harbor porpoise).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E

Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;

Odontoceti; Phocoenidae; Phocoena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAX-2001, integrated into UniProtKB/TrEMBL.
01-MAR-2001, sequence version 1.
07-FBB-2006, entry version 14.
Transferrin (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR00422; TRANSFERRIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                 The German cDNA Consortium;
                                                                                                                                                                                                                                                                                       TISSUE=Cortex;
                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9742;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GO:0005576, C:extracellular region; IEA.
GO:0008199; F:ferric iron binding; IEA.
GO:0006879; P:iron ion homeostasis; IEA.
GO:0006826; P:iron ion transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9GL95; 1-81.
        CR859370; CAH91543.1;
)5R9L7; 23-346.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF304100; AAG32042.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VVARSMGGKEDLIWELL 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VVARSMGGKEDLIWELL 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9379 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7B5DE6D524445A45 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 86;
Pred. No.
                                  mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Euteleostomi; Cetacea;
                                                                                                                                                                                                                                Amid
                                                                                                                                                                                                                                                                                                                                                                                                                                Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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RESULT 4
    REPRESENTATION OF THE PROPERTY OF THE PROPERTY
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                            Schaeffer E., Lucero M.A., Jeltsch J.-M., Py M Chambon P., Cohen G.N., Zakin M.M.; "Complete structure of the human transferrin g analogous chicken gene and human pseudogene."; Gene 56:109-116(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TRFE HUMAN STANDARD; PRT; 698 AA.

102787; 041890; 09NQBB; 09UHV0;

21-UUL-1986, integrated into UniProtKB/Swiss-Prot
01-OCT-1989, sequence version 2.

07-MAR-2006, entry version 84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
MEDLINE=88056305; PubMed=3678832; DOI=10.1016/0378-1119(87)90163-6;
Schaeffer E., Lucero M.A., Jeltsch J.-M., Py M.-C., Levin M.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
NUCLEOTIDE SI
                                                                                                                   Beutler E., Gel
Fairbanks V.F.;
                                                                                                                                                                NUCLEOTIDE SEQUENCE, AND VARIANT ATRANSFERRINEMIA MEDLINE=20563920; PubMed=11110675;
                                                                                                                                                                                                                                       Hershberger C.L., Larson J.L., Arnold B., Rosi
Williams P., Dehoff B., Dunn P., O'Neal K.L.,
"A cloned gene for human transferrin.",
Ann. N. Y. Acad. Sci. 646:140-154(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yang F., Lum J.B., McGill J.R., Moore C.M. van Bragt P.H., Baldwin W.D., Bowman B.H.; "Human transferrin: cDNA characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  binding globulin).
Name=TF; ORFNames=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Serotransferrin precursor (Transferrin)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical
SEQUENCE 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN_1;

PROSITE; PS00206; TRANSFERRIN_2;

PROSITE; PS00207; TRANSFERRIN_3;
                                                                          Blood
                                                                                                                                                                                                                                                                                                                                MEDLINE=92231399; PubMed=1809186;
                                                                                                                                                                                                                                                                                                                                                              TISSUE-L:
                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=84194084; PubMed=6585826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00405; Transferrin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                localization."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         270
                                                                          96:4071-4074 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VVARSMGGKEDLIWELL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORFNames=PRO1400;
  SEQUENCE [GENOMIC DNA]
Carrington D.P., da
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE,
                                                                                             characterization
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 protein.
697 AA; 7
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                                                                                                                                            Gelbart
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Euarchontoglires; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77010 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AND VARIANTS TF*B2;
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                                                                                                                                               Lee
                                                                                                                                          P.L.,
                                                                                             of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 86;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       81:2752-2756 (1984)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Moore C.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
  Ponte S.H.,
                                                                                                case
                                                                                                                                               Trevino
                                                                                             ō,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vertebrata; Euteleostomi;
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                                                                                                atransferrinemia.";
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                                                                                                                                                                                                                                                                                                          Rosteck
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Naylor S.L.,
  Hastings N.C.,
                                                                                                                                               Fernandez M.A.,
                                                                                                                                                                                                                                                                                      Riemen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Catarrhini;
                                                                                                                                                                                                                                                                                 P.R. Jr.,
                                                                                                                                                                                              PRO-477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Comparison with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Beta-1-metal-
                                                                                                                                                                                                                                                                                           Tice
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buertow, K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buertow, K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Souldin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., McCEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Scheutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schentz J., Myers R.M.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalaka U., Smailus D.E.,
Tend mouse converse C.M., Marra M.A.;
Tend mouse for Marra M.A.;
                                                                                                                                                                                                                                                                                                              McGillivray R.T.A., Men
Lineback-Zins J., Brew
"The primary structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE OF 422-698.
MEDLINE=84153910; PubMed=6322780;
MEDLINE=84153910; Park I., Besmo
                                                                                                                                                                      "Organization of the human transferrin originated by gene duplication.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Functional prediction of the coding
by analysis of cDNA clones from humar
Submitted (JAN-1999) to the EMBL/GenE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Fetal
Zhang C., Yu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and
                                                                                                                                                                                                                                                                                      structure."
                                                                                                                                                                                                                                                                                                                                                                                                                              Zakin M.M., Kahn A.; "Molecular cloning and sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "SeattleSNPs. NHLBI HL66682 FHCRC, Seattle, WA (URL: htt
              MEDLINE=87192006;
                                                                                                              MEDLINE=87066744;
                                                                                                                                                                                                                               MEDLINE=85216459;
                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                              MEDLINE=83160878;
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                                                                                                                                                                                                                                                                                                                                                                                                                      transferrin."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=22388257;
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 Adrian
                             NUCLEOTIDE
                                                                                                                                NUCLEOTIDE SEQUENCE OF 1-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mouse cDNA sequences.";
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                                                                                                  M. A.
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ted (MAY-2003)
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                                                                                    region of the
                                                                                                                                                                                                                  Schaeffer
                                                                                                                                                                                                                                                                         Chem.
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                             SEQUENCE OF 1-72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           liver;
Y., Zh
                                                                                                S744; PubMed=3786138;
Schaeffer E., Cohen G.N.,
 Korinek B.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kuldanek S.A.,
                                                                                                                                                                                                                                                                                                     bromide
                                                                                                                                                                                                                                                                         258:3543-3553 (1983)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; PubMed=12477932; DOI=10.1073/pnas.242603899;
Feingold E.A., Grouse L.H., Derge J.G.,
                                                                                                                                                            Sci.
                                                                                                                                                                                                                                                                                                                                                            PubMed=6833213;
                                                                                                                                                                                                               PubMed=3858812;
r E., Sidoli A.,
             PubMed=3106157;
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                                                          14:8692-8692(1986)
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http://pga.gs.washington.edu).";
the EMBL/GenBank/DDBJ databases.
 Bowman B.H.,
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             DOI=10.1016/0378-1119(86)90277-5;
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X-RAY CRYSTALLOGRAPHY (1.6 ANGSTROMS) OF 22-350.
MEDLINE=98272665; PubMed=9609685; DOI=10.1021/bi980355j;
Macgillivray R.T.A., Moore S.A., Chen J., Anderson B.F.,
Macgillivray R.T.A., Moore S.A., Murphy M.E.P., Wang Y.,
Luo Y., Bewley M.C., Smith C.A., Murphy M.E.P., Wang Y.,
Woodworth R.C., Brayer G.D., Baker E.N.;
"Two high resolution crystal structures of the recombinary."
                                                                                                                                                                                                                                                                Woodworth R.C., Mason A.B., Funk W.D., Mc "Expression and initial characterization of the N-terminal half-molecule of human Biochemistry 30:10824-10829 (1991).
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MEDLINE=20392111; PubMed=10931525;

DOI=10.1002/1097-4547(20000815)61:44.388::AID-JNR5>3.0.CO;2-Q;

de Arriba Zerpa G.A., Saleh M.-C., Fernandez P.M., Guillou F.
Espinosa de los Monteros A., de Vellis J., Zakin M.M., Baron
"Alternative splicing prevents transferrin secretion during
differentiation of a human oligodendrocyte cell line.";
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Gene 49:167-175(1986).
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                                                                                                                                                                                                          CARBOHYDRATE-LINKAGE SITES ASN-432 AND ASN-630.
PubMed=16335952; DOI=10.1021/pr0502065;
Liu T., Qian W.-J., Gritsenko M.A., Camp D.G. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease."
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Duguid J.R., Bohmont C.W., Liu N.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE OF 564-624.
Tsuchida S., Ikemoto S., Kajii E.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ
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Namekata K., Oyama F., Imagawa M., Ihara Y.;
"Human transferrin ("f): a single mutation at codon 570 dete
Biochemistry 37:7919-7928(1998)
                 release."
                            human transferrin reveal a
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McGillivray R.T.A., Mendez E., Sinha
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100:457-458(1997).
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                                               recombinant
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Matches 17
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Ensembl; ENSG0000091513; Homo sapiens.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0006199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
GO; GO:0006826; P:iron jon transport; IEA.
InterPro; IPR001156; Peptidase S60;
PANTHER; PTHR11465; Peptidase S60;
Pfam; PF00405; Transferrin; 2.
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X-RAY CRYSTALLOGRAPHY (2.2 ANGSTROMS) OF 22-350.

MEDLINE-98434369; PubMed-9760232; DOI=10.1021/bi9812064;

Jeffrey P.D., Bewley M.C., Macgillivray R.T.A., Mason A.B.,

Woodworth R.C., Baker E.N.;

"Ligand-induced conformational change in transferrins: crystal

structure of the open form of the N-terminal half-molecule of
NON TER
                       PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1;
PROSITE; PS00206; TRANSFERRIN 2;
PROSITE; PS00207; TRANSFERRIN 3;
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07-FEB-2006, entry version 4.
Transferrin variant (Fragment).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; C:
Mammalia; Eutheria; Euarchontog
                                                                                                                                                                                                                                           EMBL; AK222755; BAD96475.1; -;
                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                  Suzuki Y., Sugano S., Totoki Y., Toyoda A., Takeda T., S
Tanaka A., Yokoyama S.;
Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=98038986; PubMed=9373149; DOI=10.1016/S0378-1119(97)00411-3; Suzuki Y., Yoshitomo K., Maruyama K., Suyama A., Sugano S.; "Construction and characterization of a full length-enriched and a 5-end-enriched cDNA library."; Gene 200:149-156(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Oligo-capping: a simple method to replace the eucaryotic mRNAs with oligoribonucleotides.";
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Q53H26;
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 A54775D23B9A4FFF CRC64;
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01-MAR-2001, integrated into UniProtKB/TrEMBL.
01-MAR-2001, sequence version 1.
07-FEB-2006, entry version 14.
Transferrin (Fragment).
Mesoplodon peruvianus (Peruvian beaked whale).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
Odontoceti; Ziphiidae; Mesoplodon.
NCBI_TaxID=27617;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ziphius cavirostīis (Goose-beaked whale).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea;
Odontoceti; Ziphiidae; Ziphius.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR001156; Peptidase S60.
PANTHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HSSP; P02787; 1A8E.

SMR; Q9GL91; 1-80.

GO; GO:0005576; C:extracellular region; IE

GO; GO:0008199; F:ferric iron binding; IEA

GO; GO:0006879; P:iron ion homeostasis; IE

GO; GO:0006879; P:iron ion transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                      Q9GL92_MESPE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81 AA;
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Pred. No.
                                                                                       Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 81;
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PRELIMINARY; PR
Q3YAS1;
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27-SEP-2005, integrated into Uni
27-SEP-2005, sequence version 1
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SMART; SM
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NUCLEOTIDE SEQUENCE.

MEDLINE=20481910; PubMed=11027333; DOI=10.1073/pnas.97.21.11343; Cassens I., Vicario S., Waddell V.G., Balchowsky H., Van Belle D. Ding W., Fan C., Mohan L., Simoes-Lopes P.C., Bastida R., Meyer J. Stanhope M.J., Milinkovitch M.C.;

"Independent adaptation to riverine habitats allowed survival of ancient cetacean lineages.";

Proc. Natl. Acad. Sci. U.S.A. 97:11343-11347(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; V
Mammalla; Eutheria; Euarchontoglires; Pri
Cercopithecidae; Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GG; GG:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GG:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase S60.
PANTHER; PTHR111485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 1.
                                                                                                                                                                      EMBL; DQ147929; AAZ80920.1; -; mRNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron bing; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase S60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cells in primate experimental myopia."; Submitted (JUL-2005) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Form deprivation modulates proliferation of retinal cells in primate experimental myopia.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; I.
PROSITE; PS00206; TRANSFERRIN_2;
PROSITE; PS00207; TRANSFERRIN_3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Raviola E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
Tkatchenko A.V., Walsh P.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Transferrin (Fragment).
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                                                                                                                                                          Pfam; PF00405; Transferrin;
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Q9GL92; 1-81.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1DC9418BF5FEACC3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tkatchenko T.V.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            see http://www.uniprot.org/terms
Attribution-NoDerivs License
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RESULT 10
Q9GL94 INIGE
ID Q9GL94 INIGE
AC Q9GL94;
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QARSMI_MACFA
ID QARSMI_M
AC QARSMI_
DT 19-UUL-2
DT ACACACA E
COC CETCOPIT
COC MAMMMALIA
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RX PUBMEd=1
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RY POSITE;
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DR GO; 
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Best Local S
Matches 16
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Q4R5M1;
19-JUL-2005;
19-JUL-2005;
19-JUL-2006;
07-FEB-2006;
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Brain cDNA, clone: QccE-13766, similar to human transferrin Macaca fascicularis (Crab eating macaque) (Cynomolgus monke Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo Mammalia; Eutheria; Buarchontoglires; Primates; Catarrhini; Cercopithecidae; Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR001156; Peptidase_S60. 1. PANTHER; PTHR11485; Peptidase_S60; 1. Pfam; PF00405; Transferrin; 2. PRINTS; PR00422; TRANSFERRIN. SMART; SM00094; TR_PER; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GO; GÖ:0005576; C:extracellular region; IEA. GO; GO:0008199; F:ferric iron binding; IEA. GO; GO:0006879; P:iron ion homeostasis; IEA. GO; GO:0006826; P:iron ion transport; IEA. InterPro; IPR001156; Peptidase_S60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=15944441; DOI=10.1093/molbev/msil07;
Osada N., Hirata M., Tanuma R., Kusuda J., Hida M., Suzuki Y.,
Sugano S., Gojobori T., Shen C.-K.J., Wu C.I., Hashimoto K.;
Sugano S., Gojobori T., Shen C.-K.J., Wu C.I., Hashimoto K.;
"Substitution Rate and Structural Divergence of 5'UTR Evolution:
Comparative Analysis Between Human and Cynomolgus Monkey cDNAs."
Mol. Biol. Evol. 22:1976-1982(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               International consortium for macaque cDNA sequipun sequences of macaque genes expressed in revolutionary implications.";
Submitted (JUN-2005) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS00205; TRANSFERRIN_1; 2.
PROSITE; PS00206; TRANSFERRIN_3; 2.
PROSITE; PS00207; TRANSFERRIN_3; 2.
SEQUENCE 698 AA; 77010 MW; F19A15A7C898399B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; AB169522; BAE01604.1; -; mRNA. Q4R5M1; 23-346.
                                                                                                                                                                 270
                                                                                                                                                                                                                                                                 l Similarity
16; Conserv
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nilarity 94.1%;
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                                PRELIMINARY;
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153
17082 MW;
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Pred. No. 1.2e-05;
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Pred. No. 6.3e-05;
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Best Local S
Matches 16
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9GCL93 PHYCA PRELIMINARY; PRT; 81 AA.
09GL93,
01-MAR-2001, integrated into UniProtKB/TrEMBL.
01-MAR-2001, sequence version 1.
07-FEB-2006, entry version 15.
                                                                                                                                 NUCLEOTIDE SEQUENCE.

MEDLINE=20481910; PubMed=11027333; DOI=10.1073/pnas.97.21.11343;
Cassens I., Vicario S., Waddell V.G., Balchowsky H., Van Belle D.
Ding W., Fan C., Mohan L., Simoes-Lopes P.C., Bastida R., Meyer F.
Stanhope M.J., Milinkovitch M.C.;
"Independent adaptation to riverine habitats allowed survival of ancient cetacean lineages.";
Proc. Natl. Acad. Sci. U.S.A. 97:11343-11347(2000).
                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea; Odontoceti; Physeteridae; Physeter.
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SEQUENCE
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                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transferrin (Fragment).
Physeter catodon (Sperm whale) (Physeter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/tabletributed under the Creative Commons Attribution-NoDerivs License
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MEDLLNB=20481910; PubMed=11027333; DOI=10.1073/pnas.97.21.11343;
Cassens I., Vicario S., Waddell V.G., Balchowsky H., Van Belle D.,
Ding W., Fan C., Mohan L., Simces-Lopes P.C., Bastida R., Meyer A.
Stanhope M.J., Milinkovitch M.C.;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea;
Odontoceti; Iniidae; Inia.
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01-MAR-2001, sequence version 1.
07-FBB-2006, entry version 15.
Transferrin (Fragment).
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     EMBL;
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PANTHER; PTHR11485; Peptidase_S60; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , Q9GL94; 1-81.
GG:0005576; C:sxtracellular region; IEA.
GO:000819; F:ferric iron binding; IEA.
GO:0006879; P:iron ion homeostasis; IEA.
GO:0006826; P:iron ion transport; IEA.
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P19134; lJNF.
     AF304102;
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     AAG32044.1;
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Pred. No. 1.4e-05;
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RESULT 12
TRFE_BOVIN
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SMR; Q9GL93; 1-81.

GO; GO:0005576; C:extracellular region; IEA.

GO; GO:0008199; F:ferric iron binding; IEA.

GO; GO:0006879; P:iron ion homeostasis; IEA.

R GO; GO:0006826; P:iron ion transport; IEA.

R InterPro; IPR001156; Peptidase_S60.

R PANTHER; PTHR11485; Peptidase_S60; 1.
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Best Local
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                                                                                                                                                                                                                                                                                         --- FUNCTION: Transferring are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate. It is responsible for the transport of iron from sites of absorption and heme degradation to those of storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.

--- SUBCILIULAR LOCATION: Secreted protein.

--- SUBCILIULAR LOCATION: Secreted by the liver and secreted in plasma.
--- TISSUE SPECIFICITY: Expressed by the liver and secreted in plasma.
--- MISCELLANEOUS: The peptidase $60 domains do not have protease function as the catalytic residues are missing.
--- SIMILARITY: Belongs to the transferrin family.
--- SIMILARITY: Contains 2 peptidase $60 domains.
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NON_TER
SEQUENCE
                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=96132861; PubMed=8557646; DOI=10.1074/jbc.271.2.1166; Retzer M.D., Kabani A., Button L.L., Yu R.H., Schryvers A.B.; "Production and characterization of chimeric transferring for the determination of the binding domains for bacterial transferring the state of the binding domains for bacterial transferring determination 
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01-NOV-1997, sequence versi
07-MAR-2006, entry version
                                            InterPro; IPR001156; Peptidase S60; PANTHER; PTHR11485; Peptidase S60; Pfam; PF00405; Transferrin; 2.
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Distributed
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Pecora; Bovidae; Bo
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Name=TF;
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PRINTS; PR00422; TRANSFERRIN.
PROSITE; PS00207; TRANSFERRIN_3; 1.
PRINTS; PR00422
SMART; SM00094;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biol. Chem. 271:1166-1173(1996).
                                                                                                                                                     P09571;
                                                                                                                                                                           U02564; AAA96735.1;
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Attribution-NoDerivs License
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                TREE RABIT STANDARD,
P19134; O46514;
O1-NOV-1990, integrated into UniProtKB/Swiss-Prot.
15-JUL-1999, sequence version 4.
07-MAR-2006, entry version 61.
07-MAR-2006 entry version file (Siderophical Constraint)
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TRANSFERRIN_3; 1.
transport; Iron; Iron transport;
at; Signal; Transport.
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similarity).
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Eukaryota; Metazoa; (Mammalia; Eutheria; I Oryctolagus. NCBI TaxID=9986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=91274362; PubMed=2054387; DOI=10.1016/0167-4781(91)90021-D; Banfield D.K., Chow B.K.-C., Funk W.D., Robertson K.A., Umelas T.M. Woodworth R.C., Macgillivray R.T.A.; "The nuclectide sequence of rabbit liver transferrin cDNA."; Biochim. Biophys. Acta 1089:262-265(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           X-RAY CRYSTALLOGRAPHY (2.3 ANGSTROMS).
Sarra R., Garratt R.C., Gorinsky B., Jhoti H., Lindley P.F.;
"High-resolution X-ray studies on rabbit serum transferrin:
preliminary structure analysis of the N-terminal half-molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Evans R.W., Aitken A., Patel K.J.;
"Evidence for a single glycan moiety
location of the glycan within the pol
PEBS Lett. 238:39-42(1988).
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MEDLINE-89005676; PubMed=3169252; DOI=10.1016/0014-5793(88)80221-7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Godovac-Zimmermann
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MEDLINE=88209278;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         resolution.";
cta Crystallogr. B 46:763-771(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -RAY CRYSTALLOGRAPHY (3.3
                                                                                                                                                                                                                                                     SUBUNIT: Monomer.
SUBCELLULAR LOCATION: Secreted protein.
TISSUE SPECIFICITY: Expressed by the liver and a miscellaneous: The peptidase S60 domains do not function as the catalytic residues are missing.
SIMILARITY: Belongs to the transferrin family.
SIMILARITY: Contains 2 peptidase S60 domains.
                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: Transferrins are iron binding transport proteins we can bind two Fe(3+) ions in association with the binding of anion, usually bicarbonate. It is responsible for the transpiron from sites of absorption and heme degradation to those storage and utilization. Serum transferrin may also have a frole in stimulating cell proliferation.
    X58533; CAA41424.1;

AF031625; AAB94136.1;

AF031611; AAB94136.1;

AF031612; AAB94136.1;

AF031613; AAB94136.1;

AF031614; AAB94136.1;

AF031615; AAB94136.1;
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A.A., Thepot D.,
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                                                                                                                                                                                           UniProt Consortium, see http://www.uniprot.org/termsthe Creative Commons Attribution-NoDerivs License
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    ; -; Genomic_DNA.; JOINED; Genomic_; JOINED; Genomic_; JOINED; Genomic_; JOINED; Genomic_; JOINED; Genomic_; JOINED; Genomic_;
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., Mydin A.
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., Sarra R., Watson
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DR EMBL; AF031616; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031618; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031618; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031620; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031621; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031622; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031623; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031624; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031626; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031624; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031626; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031626; TRANSFERRIN 2: 2.

REMBL; AF031626; TRANSFERRIN 2: 2.

REMBL; AF031626; TRANSFERRIN 3: 2.

REMBL; AF031626; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF031626; AAB94136.1; JOINED; Genomic DNA.

REMBL; AF0316
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Iron 1.
Iron 1.
Iron 2.
Iron 2.
Iron 2.
Iron 2.
Iron 2.
Carbonate 1.
Carbonate 1; via amide nitrogen.
Carbonate 1; via amide nitrogen.
Carbonate 2; via amide nitrogen.
Carbonate 3; via amide nitrogen.
Carbonate 4; via amide nitrogen.
Carbonate 5; via amide nitrogen.
Carbonate 6; via amide nitrogen.
Carbonate 7; via amide nitrogen.
Carbonate 8; via amide nitrogen.
Carbonate 1; via amide nitrogen.
Carbonate 2; via amide nitrogen.
Carbonate 2; via amide nitrogen.
Carbonate 2; via amide nitrogen.
Carbonate 3; via amide nitrogen.
Carbonate 4; via amide nitrogen.
Carbonate 2; via amide nitrogen.
Carbonate 2; via amide nitrogen.
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Carbonate 2; via amide nitrogen.
Carbonate 3; via amide nitrogen.
Carbonate 4; via amide nitrogen.
Carbonate 5; via amide nitrogen.
Carbonate 6; via amide nitrogen.
Carbonate 6; via amide nitrogen.
Carbonate 6; via amide nitrogen.
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/FTId=PRO 0000035717.
Peptidase S60 1.
Peptidase S60 2.
Iron 1.
Iron 1.
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(in Ref. 3
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RESULT 14

ICA PIG

ID ICA PIG

ID ICA PIG

AC Q29545;

DT 15-JUL-1999, integrated into UniPr

DT 01-NOV-1996, sequence version 1.

DT 07-FEB-2006, entry version 39.

DE Inhibitor of carbonic anhydrase pr

GN Name=ICA;

OS Sus scrofa (Pig).

OC Eukaryota; Metazoa; Chordata; Cran

OC Mammalia; Eutheria; Laurasiatheria

OC Sus

NCBI_TaxID=9823;

RN [1]

RP NUCLEOTIDE SEQUENCE [MRNA], AND PA

MEDLINE=97254619; PubMed=9100029;

RA Wuebbens M.W., Roush E.D., Decastr
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                              270
                                                                                                                                                                                                                                                                                                                                                                        1 VVARSWGGKEDLIWELL 17
|||||:||||||||
70 VVARSVDGKEDLIWELL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ch 87.
| Similarity 88.
| 15; Conservative
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Laurasiatheria; Cetartiodactyla; Suina; Suida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Score 75; DB; Pred. No. 0.00
                                                                                                                                                                                                                                               UniProtKB/Swiss-Prot.
                                                                                                                                                                                       precursor
                                                                                                                                                                                                                                                                                   704 AA.
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RESULT 15
Q7TSX8_MARMO
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Best Local
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PANTHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 2.
PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR PER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 2; 1.
PROSITE; PS00207; TRANSFERRIN 3; 2.
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CARBOLFID
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-i- FUNCTION: Specifically binds and inhibits carbonic anhydrase with nanomolar affinity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HSSP; P09571; 1H7
LinkHub; Q29545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; U36916; AAB58956.1; -; mRNA.
PIR; I47228; I47228.
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Distributed
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                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBUNIT: Monomer.
SUBCELULAR LOCATION: Secreted protein.
TISSUE SPECIFICITY: Plasma.
TISSUE SPECIFICATE: Plasma.
PTM: N-glycosylated.
MISCELLANEOUS: The peptidase $60 domains do not function as the catalytic residues are missing.
SIMILARITY: Belongs to the transferrin family.
SIMILARITY: Contains 2 peptidase $60 domains.
                                                                                                             270
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                                                                                                       VVARSVDGKEDLIWELL
                                                                                                                                           VVARSMGGKEDLIWELL
                                                                                                                                                                                                                                                                                                                  by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                  Conservative
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383
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188
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88.2%;
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                                                                                                                                                                                                                                                                                                                       MW.
                                                                                                                                                                                                                                                                                                                    Potential.
Inhibitor of carbonic anhydrase.
Inhibitor of carbonic 
                                                                                                                                                           17
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                                                                                                                                                                                                          1;
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Pred. No.
                                                                                                                                                                                                               Mismatches
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Q7TSX8;
01-OCT-2003,
01-OCT-2003,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PRO0422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 3; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AY288100; AAP37129.1; -; mRNA.
HSSP; P19134; JUNF.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Sciuridae; Xerinae; Marmotini; Marmota.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
Pfam; PF00405; Transferrin; 2.
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rinaudo J.A.S., Gerin J.L.;
Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE
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l Similarity 82.4%;
14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     76466 MW;
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Pred. No. 0.0033
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GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd

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Result
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Maximum DB seq length: 200000000
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Maximum Match 10
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1: /EMC Celerra SIDS:

2: /EMC Celerra SIDS:

3: /EMC Celerra SIDS:

4: /EMC Celerra SIDS:

5: /EMC Celerra SIDS:

6: /EMC Celerra SIDS:

7: /EMC Celerra SIDS:
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/EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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US-09-037-188-2
US-09-037-188-3
US-09-285-310-2
US-09-753-385-2
US-09-733-310-4
US-09-037-188-4
US-09-037-188-4
US-09-133-310-4
US-09-145-681-4
US-09-723-385-4
US-08-456-106-4
US-08-456-106-4
US-09-655-77-4
US-09-633-739-4
US-09-724-864-54
US-09-724-864-54
US-09-421-632-2
US-08-655-640-4
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5, Appli
5, Appli
2, Appli
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75.6		75.6	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7
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US-08-456-106-6	US-08-453-703-6	US-08-145-681-6	PCT-US93-03614-2	US-09-633-739-2	US-09-265-577-2	US-08-476-798-4	US-08-456-108-2	US-09-158-313-4	US-08-464-167-4	US-08-456-106-2	US-08-453-703-2	US-08-461-333-4	US-08-154-019-4	US-08-250-308-2	US-08-145-681-2	US-08-476-798-2	US-09-158-313-2	US-08-464-167-2
Sequence 6,	Seguence 6,	Sequence 6,	Sequence 2,	Sequence 2,	Sequence 2,	Sequence 4,	Sequence 2,	Sequence 4,	Sequence 4,	Sequence 2,	Sequence 2,	Sequence 4,	Sequence 4,	Sequence 2,				
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RESULT 1
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US-08-175-158A-2
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D., HELLSTROM, KARL E.; ROSE, TIMOTHY M.; HELLSTROM, INGEGERD;
PURCHIO, ANTHONY F.; HU, SHIU-LOK; PENNATHUR, SRIDHAR
TITLE OF INVENTION: RECOMBINANT VIRUSES ENCODING THE HHUMAN
MELANOMA-ASSOCIATED ANTIGEN
NUMBER OF SEQUENCES: 6
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/7230
FILING DATE: 27-JAN-1987
PRIOR APPLICATION NUMBER: 827,313
APPLICATION NUMBER: 827,313
FILING DATE: 07-FEB-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO:4:
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                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/08175158A Patent No. 5986067
                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                         GENERAL INFORMATION:

APPLICANT: FUNK, Walter D.

APPLICANT: MACGILLIVRAY, ROSS T.A.

TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-

TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF

NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESS:

ADDRESSE: LAHIVE & COCKFIELD

ADDRESSE: LAHIVE & COCKFIELD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                          STREET: 60 St
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 696
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                                                                                                                                              60 State Street, suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        286
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TYPE: amino acids
TYPE: amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: protein
US-08-175-158A-2
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REFERENCE/DOCKET NUMBER: UVI-(
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 600 ---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09439740 Patent No. 6825037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 86; DB 1; Best Local Similarity 100.0%; Pred. No. 1.7e-06; Matches 17; Conservative 0; Mismatches 0;
                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 01
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: DeConti Giulio A.
REGISTRATION NUMBER: 31,503
REFERENCE/DOCKET NUMBER: UVI-005CP2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/832,029
FILING DATE: 06-FEB-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-D
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: WOODWORTH, ROBERT C.
TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/439,740 FILING DATE:
                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/175,158A FILING DATE: 28-DEC-1993 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNK, Walter D. MacGILLIVRAY, R
                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MASON, Anne B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM PC compatible
                                                                                                                                                                                                                                                                                                                                        Floppy disk
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Query Match
Best Local Similarity
Marches 17; Conserve
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US-08-470-058-2
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                                                                                                                                                                                                                                                                                                                                ZIP: 02173

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,058
FILING DATE: 06-JUN-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: TRI93-01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-661-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08470058
Patent No. 5817789
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 698 amino acids
                                                                                                                                                                                                                                                                                             TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lemontt, Jeffrey F.
TITLE OF INVENTION: Chimeric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
NUMBER OF SEQUENCES: 12
CORRESSES: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Heartlein, Michael W. APPLICANT: Lemontt, Jeffrey F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                             STRANDEDNESS:
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CITY: Lexington
STATE: MA
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646 VVARSMGGKEDLIWELL 662
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                          1 VVARSMGGKEDLIWELL 17
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                                                                         100.0%; Score 86; DB 1; ilarity 100.0%; Pred. No. 2.8e-06; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                   unknown
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US-09-037-188-2
; Sequence 2, Application US/09037188
; Patent NO. 6027921
; GENERAL INFORMATION:

RESULT 5

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Matches
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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PILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Ph.D., J.D., Jar
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 0723
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617,542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino acii
                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/037,188 FILING DATE: 02-MAR-1998 PRIOR APPLICATION DATA:
                                                                             COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                               NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                     APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 51.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Windows SOFTWARE: Factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              / Match 100.0%; Score 86; DB 2; 1
Local Similarity 100.0%; Pred. No. 2.8e-06;
hes 17; Conservative 0; Mismatches 0;
                                                                                                                                                                        STREET: 225 F
                                                                                                                                                            STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       646 VVARSMGGKEDLIWELL 662
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                                                                                                                                               ; ORGANISM: Homo sapiens
US-09-753-385-2
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CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1998-03-09
PRIOR FILING DATE: 1998-03-09
PRIOR FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-753-385-2
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Matches 17; Conservative
                                                                      Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                         SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Applic Patent No. 6858578
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE IN TRANSPORT
TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                LENGTH: 1074
TYPE: PRT
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LENGTH: 1074 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/285,310 FILING DATE: PRIOR APPLICATION DATA: APPLICATION NUMBER: 09/037,188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Fraser, Ph.D., J.D., Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07236/009002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
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646 VVARSMGGKEDLIWELL 662
                                 1 VVARSMGGKEDLIWELL 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09753385
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                                                                        Conservative
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                                                                      100.0%; Score 86; DB 2; 100.0%; Pred. No. 2.8e-06; ... Mismatches 0;
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Pred. No. 2.8e-06;
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                                                                                                       Length 1074;
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US-08-470-058-4

Sequence 4, Application US/08470058
Patent No. 5817789
GENERAL INFORMATION:
APPLICANT: Heartlein, Michael W
APPLICANT: Lemontt, Jeffrey F.

Heartlein, Michael W. Lemontt, Jeffrey F.

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; TOPOLOGY: unknown ; MOLECULE TYPE: protein US-08-470-058-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                       STATE: MA
COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 4:
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            COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS
TITLE OF INVENTION: IN TRANSPORT OF A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/470,058 FILING DATE: 06-JUN-1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
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COUNTRY: US
ZIP: 02173
                                                                                                                                COMPUTER: IBM Con
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: un
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REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: TKT93-01
                                                                                                                                                                                                                                                                                        CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
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                                                                                                                                                                                                                                                                                      Boston
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06-JUN-1995
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SELECTED SUBSTANCE INTO CELLS
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; MOLECULE TYPE: protein US-09-285-310-4
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1410 amin
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Patent No. 6262026
GENERAL INFORMATION:
APPLICANT: Heartlein, Michael W
APPLICANT: Lemontt, Jeffrey F.
Query Match 100.0%; Score 86; DB 2; Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 17; Conservative 0; Mismatches 0;
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                                                                                                                                                        TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   NAME: Fraser, Ph.D., J.D., V
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PR
TITLE OF INVENTION: IN TRANSPOR
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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CITY: Boston
STATE: MA
                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                         LENGTH:
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GY: unknown
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100.0%; Pred. No. 3.8e-06;
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IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
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                                                                                                                                                                                                                                                                                                          Janis K.
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                                   Length 1410;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 1410;
   Indels
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VVARSMGGKEDLIWELL VVARSMGGKEDLIWELL

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APPLICANT: Heartlein, Michael W.
APPLICANT: Hemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS
FILE REFERENCE: 10278-025004
CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1998-03-09
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1998-03-09
PRIOR APPLICATION NUMBER: US 09/285,310
PRIOR FILING DATE: 1999-04-02
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; ORGANISM: Homo sapiens
US-09-753-385-4
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4, Application US/08145681 Patent No. 5571691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 17; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/09753385 Patent No. 6858578
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                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: FLODPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTION PATA:
APPLICATION NUMBER: US/08/145,681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Production of Recombinant Lac
TITLE OF INVENTION: and Lactoferrin Polypeptides
TITLE OF INVENTION: Various Organsims
                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                           FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                     COUNTRY: U
ZIP: 77002
                                                                                                                                                                                                                                                                                                                      CITY: Houston
                                                                                                                                                                                                                                                                                                                                            STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           982 VVARSMGGKEDLIWELL 998
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McGregor, Martin L.
                                                                                                                                                                                                                                                                                                   Texas
                                                                                                                                                                                                                                                                                                                                            E: Baker & Botts, L.L.P.
910 Louisiana St
                                                                                                                                                                                                                                                                         USA
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                                                                                                                                    Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Using cDNA Sequences
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION UNMER: US/08/453,703
FILING DATE: Concurrently herewith
CLASSIFICATION NUMBER: US/08/453,703
FILING DATE: Concurrently herewith
CLASSIFICATION NUMBER: 08/145,681
FRIOR APPLICATION NUMBER: 08/145,681
FILING DATE: October 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8206-024
TELECOMPUNICATION: INFORMATION:
TELEPHONE: 415-854-3660
TELEPAX: 415-854-3660
TELEPAX: 415-854-3660
TELEPAX: 415-854-3660
TELEPAX: 415-854-3660
TELEPAX: 110-06161
SEQUENCE CHARACTERISTICS:
LENGTH: 708 amino acids
TYPE: amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: May, Gregory S.

TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using CDI
TITLE OF INVENTION: Various Organisms
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: pro
HYPOTHETICAL: YES
ANTI-SENSE: NO
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/229/1874
MOLECULE TYPE: protein HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Conneely, Orla M
APPLICANT: Headon, Denis R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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CITY: New York
STATE: New York
                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10036
                                                         TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 708 amino acids
TYPE: amino acid
TOPOLOGY: linear
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1155 Avenue of the Americas
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                                          linear
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Pred. No. 0.00072;
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; ORGANISM: Bos
US-08-453-703-4
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US-08-456-106-4
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Best Local Similarity
                                                                     Matches
                                                                                Query Match
Best Local Similarity
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FILING DATE: Concurrently herewith
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,681
FILING DATE: October 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8206-025
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APPLICANT: Connec
APPLICANT: Headon
                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                               ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Bos taurus
                                                                                                                                                                                       MOLECULE TYPE: protein HYPOTHETICAL: YES ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using cDNA Sequences
TITLE OF INVENTION: Various Organisms
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ADDRESSEE: Pennie &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                         LENGTH: 708 amino acids TYPE: amino acid TOPOLOGY: linear
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CITY: New York
STATE: New York
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274 VVARSVDGKEDLIWKLL 290
274 VVÁRSVDGKEDLÍWKLL 290
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                     1 VVARSMGGKEDLIWELL 17
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                                                                82.6%; Score 71; Initiative 82.4%; Pred. No. 0. Conservative 2; Mismatche
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82.4%; Pred. No. 0.00072;
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                                                                    Mismatches
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                                                                                                  Length 708;
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Search completed: June 13, 2006, 04:25:49 Job time: 30 secs
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                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/456,10:
APPLICATION UMBER: US/08/456,10:
FILING DATE: CONCURRENTLY herewii
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,681
FILING DATE: OCCODER 28, 1993
ATTONNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 2506-02
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
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APPLICANT: Connee
APPLICANT: Headon
                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 415-854-3694
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                  HYPOTHETICAL: YE ANTI-SENSE: NO ORIGINAL SOURCE: ORGANISM: Bos
                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 708 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
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TITLE OF INVENTION: Produc
TITLE OF INVENTION: and La
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                  Local
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                                                                 274 VVARSVDGKEDLIWKLL 290
                                                                                1 VVARSMGGKEDLIWELL 17
                                                                                                                                  14;
                                                                                                                                                  Similarity
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MATion.
Conneely, Original R.
Thadon, Denis R.
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                                                                                                                                  Conservative
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                                                                                                                                                                                                                   Bos taurus
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                                                                                                                                                82.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Production of Recombinant Lactoferrin
and Lactoferrin Polypeptides Using cDNA Sequences in
Various Organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Orla M.
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                                                                                                                                                  Score 71; DB 2;
Pred. No. 0.00072;
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                                                                                                                                                                 Length 708;
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                                                                                                                                  0;
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Result
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Maximum Match 100%
Listing first 45 s
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length: 2000000000
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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    GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.
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                   US-10-378-094-2

US-10-384-060-2

US-10-314-94-2

US-10-383-201-10

US-10-383-201-14

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US-10-387-711-2

US-10-513-523-3

US-11-177-506-46
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US-10-429-653-4

US-10-429-659-4

US-10-429-661-4

US-10-429-660-4

US-10-429-662-4

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US-10-384-060-3

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66	66	66	66	66	66	66	66	66	69	71	71	71	76	86	86	86	86
76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	76.7	80.2	82.6	82.6	82.6	88.4	100.0	100.0	100.0	100.0
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US-10-316-253-275	US-10-316-253-273	US-10-023-096-2	US-10-639-835-4	US-10-077-381-4	US-10-076-816-4	US-10-169-297-49	US-10-408-765A-241	US-09-864-408A-5324	US-10-801-990-183	US-10-513-523-2	US-10-620-256-4	US-10-169-297-8	US-10-513-523-4	US-10-473-127-804	US-10-473-127-805	US-09-753-385-4	US-09-753-385-2
Sequence 275, App	Sequence 273, App	Sequence 2, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 49, Appl	Sequence 241, App	Sequence 5324, Ap	Sequence 183, App	Sequence 2, Appli	Sequence 4, Appli	Sequence 8, Appli	Sequence 4, Appli	Sequence 804, App	Sequence 805, App	•	Sequence 2, Appli

; LENGTH: 17 ; TYPE: PRT ; ORGANISM: Homo s US-10-612-162-1 APPLICANT: Dade Behring Marburg GmbH TITLE OF INVENTION: Carbohydrate deficient transferrir TITLE OF INVENTION: antibodies, their preparation and FILE REFERENCE: 2002/B001 CURRENT APPLICATION NUMBER: US/10/612,162 CURRENT FILING DATE: 2003-07-03 PRIOR APPLICATION NUMBER: 10230550.1 PRIOR FILING DATE: 2002-07-05 PRIOR FILING DATE: 2002-07-05 PRIOR APPLICATION NUMBER: 03011334.4 PRIOR FILING DATE: 2003-05-19 NUMBER OF SEQ ID NOS: 4 SOFTWARE: Patentin Ver. 2.1 RESULT 2 US-10-384-060-76 S RESULT 1 US-10-612-162-1 ; Sequence 1, Application US/10612162 ; Publication No. US20040014145A1 ; GENERAL INFORMATION: SUFINE SEQ ID NO 1 Matches Query Match Best Local l Similarity 17; Conserv 1 VVARSMGGKEDLIWELL 17 VVARSMGGKEDLIWELL Conservative sapiens 100.0%; Score 86; 100.0%; Pred. No. <u>,</u> deficient transferrin (CDT)-specific their preparation and use Mismatches 2e-07; DB 4; Length 17; Indels 0 Gaps

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Sequence 76, Application US/10384060

Publication No. US20030226155A1

GENERAL INFORMATION:
APPLICANT: SADEGHI, Homayoun
APPLICANT: PRIOR, Christopher P.
APPLICANT: PRIOR, Christopher P.
APPLICANT: TURNER, Andrew
FILLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
FILE REFERENCE: 54710-5004-US
CURRENT APPLICATION NUMBER: US/10/384,060
CURRENT FILING DATE: 2003-03-01
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2001-08-30

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APPLICANT: LAI, Char-Huei
APPLICANT: LAI, Char-Huei
APPLICANT: LAI, Char-Huei
APPLICANT: LAI, Char-Huei
APPLICANT: SADEGHI, Homayoun
APPLICANT: TURNER, Andrew
ITILE OF INVENTION: MODIFIED TRANSFERRIN FUSION PROTEINS
FILE REFERENCE: 54710-5001-01-US
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT FILING DATE: 2003-03-04
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR PILING DATE: 2001-10-30
NUMBER OF SEG ID NOS: 66
SOFTWARE: PatentIn version 3.2
SEG ID NO 3
LENGTH: 679
TYPE: PRT
TYPE: PRT
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              Sequence 3, Application US/10384060
Publication No. US20030226155A1
GENERAL INFORMATION:
APPLICANT: SADEGHI, Homayoun
APPLICANT: TURNER, ADTENDER: P.
APPLICANT: TURNER, ADTENDER: USACREW
TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
FILE REFERENCE: 54710-5004-US
CURRENT APPLICATION UNMERR: US/10/384,060
CURRENT FILING DATE: 2003-03-10
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Publication No. US20030221201A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 80
SOFTWARE: PatentIn version 3.2
SEQ ID NO 76
LENGTH: 220
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PRIOR APPLICATION NUMBER: US 10/231,494
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NAME/KEY: MISC_FEATURE
OTHER INFORMATION: Mature Transferrin Protein
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OTHER INFORMATION: mTf sequence in prex0080
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PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 80
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; NAME/KEY: MISC_FEATURE
; OTHER INFORMATION: Mature Transferrin Protein
US-10-384-060-3
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APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The Diagnosis, Imaging and Treatment Of Tun
FILE REFERENCE: 2537.000001
CURRENT APPLICATION NUMBER: US/10/429,482
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/10231494
publication No. US20040023334A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
                                                                                                                                                              Sequence 4, Application US/10429482 Publication No. US20040219097A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3
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Best Local
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CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/315,745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
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17; Conserv
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100.0%; Pred. No. 1.1e-05;
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; LENGTH: 679 ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-429-482-4

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251 VVARSMGGKEDLIWELL 267

1 VVARSMGGKEDLIWELL 17

Matches Query Match

Local Similarity mes 17; Conserv

100.0%; Score 86; ilarity 100.0%; Pred. No. Conservative 0; Mismatch

Mismatches

1.1e-05; DB 5;

Length 679;

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Gaps

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US-10-429-497-4 RESULT 7

Sequence 4, Application US/10429497
Publication No. US20040219098A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For The Treatment of Tumors
FILE REFERENCE: 2537.00006
CURRENT APPLICATION NUMBER: US/10/429,497
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1

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Sequence 4, Application US/10429515

Publication No. US20040219099A1

GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir

ITILE OF INVENTION: Methods For The Treatment Of Tumors

FILE REFERENCE: 2537.000005

CURRENT APPLICATION NUMBER: US/10/429,515

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 6

SOUTWARE: Patentin version 3.1

SEQ ID NO 4

LENGTH: 679

TYPE: PRT
ORGANISM: Homo sapiens

US-10-429-515-4
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; LENGTH: 679
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-429-497-4
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US-10-429-598-4
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GENERAL INFORMATION:
             Sequence 4, Application US/10429598
Publication No. US20040219100A1
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Best Local Similarity 100.0%; Pred. No. 1.1e-05;
Matches 17; Conservative 0; Mismatches 0;
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; TYPE: PRT ; ORGANISM: Homo sapiens US-10-429-598-4
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US-10-429-653-4
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US-10-429-635-4
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US-10-429-635-4
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LENGTH: 679
TYPE: PRT
                                                                                                                                                                                           APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Compositions For Drug Delivery
FILE REFERENCE: 2537,00009
CURRENT APPLICATION NUMBER: US/10/429,653
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
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Publication No. US20040219101A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Composition Useful For The Treatment of Tumors
FILE REFERENCE: 2537.000003
CURRENT APPLICATION NUMBER: US/10/429,598
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
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Publication No. US20040219102A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Composition Useful For Treatment Of Tumors
FILE REFERENCE: 2537.000007
CURRENT APPLICATION NUMBER: US/10/429,635
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.1
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TYPE: PRT
ORGANISM: Homo sapiens
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Local Similarity 100.0%; Pred. No. 1.1e-05;
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Local Similarity 100.0%;
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Pred. No. 1.1e-05;
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US-10-429-659-4
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                                   NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 679
                                                                                                          Sequence 4, Application US/10429660
Publication No. US20040220084A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods for Nucleic Acid Delivery
FILE REFERENCE: 2537.000012
CURRENT APPLICATION NUMBER: US/10/429,660
CURRENT FILING DATE: 2003-05-02
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LENGTH: 679
TYPE: PRT
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Publication No. US20040219104A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For Treatment Of
FILE REFERENCE: 2537.000008
CURRENT APPLICATION NUMBER: US/10/429,661
CURRENT FILING DATE: 2003-05-02
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Matches
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APPLICANT: Sandhu, Jasbir
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods Useful For The Diagnosis, Imaging and Treatment Of Tumors
FILE REFERENCE: 2537.000004
CURRENT APPLICATION NUMBER: US/10/429,659
CURRENT FILING DATE: 2003-04-30
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Best Local (
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
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Publication No. US20040219103A1
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SOFTWARE: PatentIn version 3.1
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TYPE: PRT
ORGANISM: Homo sapiens
TYPE: PRT
ORGANISM: Homo sapiens
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Sequence 4, Application US/10429662

Publication No. US20040220085A1

GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Compositions For Nucleic Acid Delivery
FILE REFERENCE: 2537.000010

CURRENT APPLICATION NUMBER: US/10/429,662

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 12

SOFTWARE: PatentIn version 3.1

SEQ ID NO 4

LENGTH: 679

TYPE: PAT

ORGANISM: Homo sapiens
Search completed: June 13, Job time: 98.75 secs
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US-10-429-662-4
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Best Local Similarity 100.0%;
Matches 17; Conservative 0
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Pred. No. 1.1e-05;
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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                                     US-10-515-429-37

US-11-258-767-23

US-11-258-767-13

US-11-258-767-16

US-11-258-767-19

US-11-258-767-29

US-11-258-767-29

US-11-258-767-39

US-11-258-767-39
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45	44	43	42	41	40	9	38	37	36	35	34	LJ LJ	32	31	30	29	28	27	26
38	40	40				41	41	55	55	60	65	65	66	66	66	66	66	66	66
44.2	46.5	46.5	46.5	47.7	47.7	47.7	47.7	64.0	•	69.8	•	75.6	76.7	76.7	76.7	76.7	76.7	76.7	76.7
146	206	177	176	654	634	601	581	707	707	686	703	696	711	711	711	711	711	711	711
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US-11-055-093-225	US-10-953-349-34871	US-10-953-349-34872	US-10-953-349-34873	US-11-189-279-15	US-11-189-279-16	US-11-189-279-17	US-11-189-279-18	US-11-258-767-20	US-11-258-767-14	US-11-258-767-36	US-11-258-767-37	US-10-515-429-42	US-11-258-767-34	US-11-258-767-30	US-11-258-767-27	US-11-258-767-21	US-11-258-767-18	US-11-258-767-17	US-11-258-767-15
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Seguence	Sequence						
225, App	34871, A	34872, A	34873, A	15, Appl	16, Appl	17, Appl	18, Appl	20, Appl	14, Appl	36, Appl	- "	•	•	•	-	21, Appl	18, Appl	17, Appl	•

RESULT 1 US-10-515-429-63

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PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILLING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILLING DATE: 2002-08-30
PRIOR FILLING DATE: 2003-03-10
PRIOR FILLING DATE: 2003-03-10
PRIOR FILLING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 63
LENGTH: 371
TYPE: PRT
ORGANISM: artificial
                                                                                                          RESULT 2
US-10-515-429-3
                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: N-domain OTHER INFORMATION: vectors US-10-515-429-63
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               Sequence 3, Application US/10515429
Publication No. US20060105387A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 63, Application US/10515429 Publication No. US20060105387A1 GENERAL INFORMATION:
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Matches 17; Conserv
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Best Local
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APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
TILE REFERENCE: 054710-5007-WO
CURRENT EPPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                      274
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Turner, Andrew J.
Sadeghi, Homayoun
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Pred.
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FILE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-00
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-07-09
NUMBER OF 580 ID NOS: 75
SEQ ID NOS: 75
SEQ ID NOS: 75
LENGTH: 679
TYPE: PRT
RESULT 4
US-10-515-429-41
); Sequence 41, Application US/10515429
; Publication No. US20060105387A1
; GENERAL INFORMATION:
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PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR RILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-07-09
PRIOR PILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Mature transferrin protein US-10-515-429-3
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Best Local Similarity
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Best Local Similarity
Matches 17; Conserv
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SEQ ID NO 2
LENGTH: 698
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Publication No. US20060105387A1
GENERAL INFORMATION:
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APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
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                                                                                                                                                             270 VVARSMGGKEDLIWELL 286
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100.0%; Pred. No.
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7.2e-07;
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Sequence 37, Application US/10515429

Publication No. US20060105387A1

GENERAL INFORMATION:

APPLICANT: Prior, Christopher P.

APPLICANT: Turner, Andrew J.

APPLICANT: Sadeghi, Homayoun

TITLE OF INVENTION: Transferrin Fusion Protein Libraries

FILE REFERENCE: 054710-5007-WO

CURRENT APPLICATION NUMBER: US/10/515,429

CURRENT FILING DATE: 2003-08-26

PRIOR APPLICATION NUMBER: PCT/US03/26779

PRIOR APPLICATION NUMBER: US 60/406,977

PRIOR APPLICATION NUMBER: US 60/406,977

PRIOR APPLICATION NUMBER: US 10/384,060

PRIOR APPLICATION NUMBER: US 10/384,060

PRIOR FILING DATE: 2003-08-30

PRIOR FILING DATE: 2003-09-30

PRIOR FILING DATE: 2003-07-09

PRIOR FILING DATE: 2003-07-09
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TYPE: PRT
; ORGANISM: Oryctolagus cuniculus
US-10-515-429-37
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APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT APPLICATION NUMBER: PCT/US03/26779
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
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SEQ ID NO 41
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Best Local S
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PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Bos taurus
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251 VVARSVDGKEDLIWELL 267
                              1 VVARSMGGKEDLIWELL 17
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                                                                                                                             Similarity
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                                                                                                     Conservative
                                                                                                                           87.2%;
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                                                                                                                             Score 75; DB 6;
Pred. No. 4.9e-05;
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Pred. No.
                                                                                                       Mismatches
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RESULT 6
US-11-258-767-23
; Sequence 23, Application US/11258767
; Publication No. US20060094082A1

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NUMBER OF SEQ ID NOS: 67
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 23
LENGTH: 708
TYPE: PRT
ORGANISM: CAMEL
US-11-258-767-23
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TYPE: PRT
ORGANISM: CAMEL
US-11-258-767-24
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Publication No. US20060094082A1
GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
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Matches
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 24
                                                                                                                                                                      Sequence 13, Application US/11258767 Publication No. US20060094082A1 GENERAL INFORMATION:
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Best Local (
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APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P03915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
                                                                                           APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
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CURRENT FILING DATE: 2005-10-26
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APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
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les 14; Conserv
                                                                                                                                                                                                                                                                                                            274 VVARSVNGKEDLIWKLL 290
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Pred. No. 0.00016;
2; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                       Score 72; DB 7;
Pred. No. 0.00016;
2; Mismatches 1
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; ORGANISM: COW
US-11-258-767-13
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US-11-258-767-16
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; ORGANISM: WATER BUFFALO US-11-258-767-19
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APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
SEQ ID NO 16
                                                                 CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.3
SEQ ID NO 19
SEQ ID NO 19
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Matches 14; Conservative
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TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
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TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
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PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
                                    LENGTH: 70
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TYPE: PRT
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82.4%;
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Pred. No. 0.00023;
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SEQ ID NO 29
LENGTH: 708
TYPE: PRT
ORGANISM: COW
US-11-258-767-29
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; ORGANISM: COW
US-11-258-767-22
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US-11-258-767-29
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US-11-258-767-22
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Best Local S
Matches 14
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Publication No. US20060094082A1

GENERAL INFORMATION:

APPLICANT: Varadhachary, Atul

APPLICANT: Glynn, Peter

APPLICANT: Petrak, Karel

APPLICANT: Engelmayer, Jose

TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF

FILE REFERENCE: HO-P02915US1

CURRENT FILING DATE: 2005-10-26

PRIOR APPLICATION NUMBER: US 60/622,176

PRIOR FILING DATE: 2004-10-26

PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 29, Application US/11258767 Publication No. US20060094082A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 22
LENGTH: 708
                                                                                                  Query Match
Best Local Similarity
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Best Local Similarity
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APPLICANT: Petrak, Karel
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
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Local Similarity 82.4%;
les 14; Conservative
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      VVARSVDGKEDLIWKLL 290
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milarity 82.4%;
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Pred. No. 0.00024;
2; Mismatches 1
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Pred. No. 0.00024;
2; Mismatches 1
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US-11-258-767-39

Sequence 39, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:

APPLICANT: Varadhachary, Atul APPLICANT: Glynn, Peter

Glynn, Peter Petrak, Karel

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LENGTH: 708
TYPE: PRT
ORGANISM: COW
US-11-258-767-35
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; ORGANISM: COW
US-11-258-767-38
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RESULT 15
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CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.3
SEQ ID NO 35
LENGTH: 708
                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.3 SEQ ID NO 38 LENGTH: 708
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Best Local
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Best Local Similarity
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APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
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APPLICANT: Engelmayer, Jose
TITILE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
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Similarity 82.4%;
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Pred. No. 0.00024;
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Pred. No. 0.00024;
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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1: geneseqp198
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       Ads13216 Human rhe
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Adx65973 Tissue fa
Aay41871 Rheumatoi
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Aae18890 Human zsi
Abe72317 Human pro
Ady39273 Human zsi
Aef38807 E. coli L
Aaw58715 Tryptic 4
Aau78849 Scrambled
Adu3791 Human CLM
Adw68715 Chromatin
Aay24901 Peptide N
Adm96841 Tissue fa
Adm96821 Tissue fa
Adm96821 Tissue fa
                                                                                                                                                                                                                                               Adg46075 Human CDT
Adr15903 Transferr
Adr15963 Transferr
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ABG69725	AAU68379	AAB39215	AAP91103	AAP50736	AEC81921	ADT79005	AEC21066	ADU87378	ADT55054	ADJ99978	ADK52246	ADJ67240	ADK17244	ADJ82185	ADH17278	ADH17195	ADF55507	ABB76441	AAW29693	AAW29691	AAWZ9690
Abg69725	Aau68379	Aab39215	Aap91103	Aap50736	Aec81921	Adt79005	Aec21066	Adu87378	Adt55054	Adj 99978	Adk52246	Adj 67240	Adk17244	Adj 82185	Adh17278	Adh17195	Adf55507	Abb76441	Aaw29693	Aaw29691	Adw29090
Polypept	Human	Human	New immun	Plasmid	Polypept	DNA-PK	Human	Peptide	E-select	Breast	Modified	C-epsilon	Primer	Peptide		Human		Peptide	TNF recep	TNF	TIME

RESULT 1 ADG46075 antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. ADG46075; 05-JUL-2002; 2002DE-01030550. 19-MAY-2003; 2003EP-00011334. 07-JAN-2004. EP1378521-A1. Homo sapiens. Human CDT peptide fragment 25-MAR-2004 ADG46075 standard; peptide; 17 (first entry) # \$

New antibody specific for carbohydrate-deficient transferrin, useful f diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization.
specific for alcoholism, bilization.
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WPI; 2004-073743/08.

Althaus H;

(DADE-) DADE

BEHRING MARBURG GMBH.

Claim 4; SEQ ID NO 1; 21pp; German.

This invention describes a novel antibody that, in aqueous solution, blinds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSWGGKEDLIWELL, TTEDSLAKIMNOSADAMSILDGGF, SKLSMGSGLNLSEEN and YEKYLGEEVVKAV (ADG46075-ADG46079) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for

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ADM96821 AAR06889

ADM96841 ADS51653

AEF38807 AAW58715 AAU78549 ADG39400 ADG03771

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Query Match 50.0%; Score 43; DB 8; Length 17; Best Local Similarity 100.0%; Pred. No. 5.4;
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ADS13216 ID ADS1 XX AC ADS1

ADS13216 standard; peptide; 17 AA.

ADS13216;

RESULT 4

Sequence 17

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Matches
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two isoforms with different glycosylation patterns. The assay involves contacting a sample containing the protein with a proteolytic enzyme, followed by detecting the content or relative content of at least one peptide fragment produced by proteolysis. Also described is a kit for the assay method comprising the proteolytic enzyme and a substrate bound specific binding partner (S1) for at least 2 of the isoforms of the proteins. The method can be used for assaying isoforms of proteins according to their glycosylation pattern to determine their concentration or relative concentration in the sample or material (e.g. blood). The method avoids use of antibodies for distringuishing between glycosylated isoforms of the proteins. The present sequence represents a transferrin peptide fragment which is used in the exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assay for differentiating protein isoforms to determine their concentrations in sample e.g. blood, involves contacting the sample with proteolytic enzyme to produce peptide fragment by proteolysis followed by
                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an assay for a protein having
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 15; 30pp; English.
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                                                                                                                                                                                    The present invention describes an assay for a protein having at least two isoforms with different glycosylation patterns. The assay involves contacting a sample containing the protein with a protectytic enzyme, followed by detecting the content or relative content of at least one peptide fragment produced by protectysis. Also described is a kit for the assay method comprising the protectytic enzyme and a substrate bound specific binding partner (S1) for at least 2 of the isoforms of the protectins. The method can be used for assaying isoforms of proteins according to their glycosylation pattern to determine their concentration or relative concentration in the sample or material (e.g. blood). The method avoids use of antibodies for distinguishing between glycosylated method avoids use of antibodies for distinguishing between glycosylated
                                                                                                                                                                                                                                                                                                                                                                                                         Assay for differentiating protein isoforms to determine their concentrations in sample e.g. blood, involves contacting the sample with proteolytic enzyme to produce peptide fragment by proteolysis followed by
                                                                                                                Sequence 17
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                                                                                                                                                          isoforms of the proteins. The present sequence represents a transferrin peptide fragment which is used in the exemplification of the present
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                                                                                                                                            invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel isolated marker for rheumatoid arthritis selected from one of many (around 400) markers defined in the specification. Rheumatoid arthritis is a chronic inflammatory disorder of the small joints which is estimated to affect 2.1 million people in the United States alone. Current approaches to treat the disease include the use of non-steroidal antinflammatory drugs (NSAIDS), which may reduce pain, swelling and inflammation, and disease-modifying anti-rheumatic drugs (DMARDS), which act to slow the progression of the disease and avoid further joint injury. These drugs are associated with a number of serious side effects and the search for improved therapeutics is a subject of active research. The marker of the invention demonstrates antiarthritic activity and may be useful in preparing a composition for diagnosing or treating rheumatoid arthritis. The current sequence is that of a human rheumatoid arthritis marker peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated marker for rheumatoid arthritis, useful in preparing composition for diagnosing or treating rheumatoid arthritis.
   23-JUL-2004; 2004WO-US023908.
                                                            10-FEB-2005.
                                                                                                                     WO2005011576-A2
                                                                                                                                                                                                                                  protein purification; chromatography; screening
                                                                                                                                                                                                                                                                                              Human serum albumin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; --
                                                                                                                                                                                                                                                                                           (HSA) peptide fragment, SEQ ID 238.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50.0%; Score 43; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schulman
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RESULT 6
ADE65973
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method for analyzing a mixture of proteins. The method involves selecting sets of peptides from several peptides, sequencing one or more portions of the peptides in two or more sets of peptides in two or more sets of peptides, to produce a data set, and determining the likelihood of the presence of two or more proteins in the mixture of proteins based on two or more data sets. The invention further comprises an article of manufacture comprising a computer-readable media with computer-readable instructions for carrying out the method. The method is useful for analyzing a mixture of proteins. The method is useful for scheroling and characterization of protein mixtures. This sequence represents a human serum abbumin (HSA) peptide fragment used in the novel protein analysis method of the invention.
             A composition for diagnosing, preventing and treating cancer (e. prostatic, renal or uterine cancer) comprises 161P2F10B polynucl
                                                                                                                                                         07-NOV-2001; 2001US-00005480
31-JAN-2002; 2002US-00062109
and polypeptides.
                                                        WPI; 2003-441560/41.
                                                                                   Jakobovits A, Morrison RK,
                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE65973 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analyzing mixture of proteins by sequencing portions of peptides in two or more sets of peptides from several peptides, to produce data set, determining likelihood of presence of proteins in mixture of proteins
                                                                                                                                                                                                                               15-MAY-2003.
                                                                                                                                                                                                                                                           WO2003040340-A2
                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                  161P2F10B; cancer;
                                                                                                                                                                                                                                                                                                                                           Human 161P2F10B protein variant 3 peptide SeqID85.
                                                                                                                                                                                                                                                                                                                                                                       29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                     ADE65973;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2005-162781/17.
                                                                                                                                                                                                 07-NOV-2002; 2002WO-US036002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; SEQ ID NO 238; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   based on data sets.
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                                                                                                                              (AGEN-) AGENSYS INC.
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                                                                                   Raitano AB,
Challita-Eid
                                                                                                                                                                                                                                                                                                                cytostatic; gene therapy; vaccine; human
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                                                                                    Faris M,
PM;
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Pred. No.
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polynucleotides
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel composition which comprises a substance that modulates the status of a novel protein (1612F10B) and its variants having a sequence of 875 amino acids provided in the specification. The protein of the invention is over-expressed in certain cancers. The compounds of the invention may have cytostatic activity and the sequence of the 1612F10B protein, and the gene which encodes it, may be useful for gene therapy or the development of a vaccine. The composition and methods of the invention are useful in diagnosing, preventing and treating cancer. The present sequence is the amino acid sequence of peptide derived from the human 161P2F10B variant 3 protein and used in the exemplification of the invention.
          The invention relates to peptide antagonists of tissue Factor VIIa (tFVIIa) and a method of inhibiting FVIIa activity by contacting FVIIa with the peptide in the presence of tissue factor and under conditions that allow binding of the compound to FVIIa to occur. The peptides are useful for preventing or treating chronic thromboembolic diseases or disorders associated with fibrin formation including vascular disorders, such as deep venous thrombosis, arterial thrombosis, stroke, atherosclerosis, or septicaemia. The present sequence represents a tFVIIa peptide antagonist of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue Factor VIIa; tFVIIa; chronic thromboembolic disease; fibrin formation; vascular disorders; deep venous thrombosis; arterial thrombosis; stroke; atherosclerosis; septicaemia.
                                                                                                                                                                                                                                      New peptide, useful for preventing or treating chronic thromboembolic diseases or disorders associated with fibrin formation including vascular disorders, such as deep venous thrombosis, arterial thrombosis, and
                                                                                                                                                                                                                                                                                                             WPI; 2004-356247/33.
                                                                                                                                                                                                                                                                                                                                            Lazarus RA, Maun HR;
                                                                                                                                                                                                                                                                                                                                                                                                               06-FEB-2002; 2002US-0355420P
                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2003; 2003US-00356257
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                                                                                                                                                                                       SEQ ID NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (tFVIIa)
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                                                                                                                                                                                      295; 102pp; English.
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Pred. No. 8.2e+02;
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CC dimensional array of features. The method can be used for screening, cd dimensional array of features. The method can be used for screening, cd diagnosis and prognosis of RA in a subject or for monitoring the effect cof an anti-RA drug or therapy administered to a subject. The method comprises: (a) analysing a sample of serum or plasma and optionally comprises: (a) analysing a sample of serum or plasma and optionally comprises: (a) analysing a sample of serum or plasma and optionally comprises; (b) identifying at least one chosen comprises and (c) comparing the abundance of each chosen feature in the sample with the abundance of that chosen feature in the presence or sample with the abundance of that chosen feature in the sample of the chosen feature in the subject. The method can also be used in clinical studies for testing drugs for therapy of RA, for purification of RA-compositic protein isoforms (RPIs), and for production of antibodies to RPIs. The RA-diagnostic feature (RAPIs), and for production of antibodies to compounds that promote or inhibit their activity, which are then used as CC RA drugs. Nucleic acid encoding RADFs can be used in gene therapy compounds that promote or inhibit their activity, which are then used as CC RAY42103 represent expression reference protein isoform peptides and CC LAR425066 to AR225068 represent degenerate probes for RPIs, which are all cc used in the exemplification of the present invention
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                           A method has been developed for the diagnosis of human rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 18; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Diagnosis of human rheumatoid arthritis by two-dimensional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Parekh RB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (OXFO-) OXFORD GLYCOSCIENCES UK LTD
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Pred. No. 1.
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Matches Best Local Query Match

Similarity 5; Conserv

Conservative

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Mismatches

1.8e+03; hes 0; DB 2;

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Gaps

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Length 17; Indels

32.6%;

Score 28; Pred. No.

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RESULT 9
AR23945
AD AR23945
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CC nucleic acids and methods for modulating gastric contractility, nutrient CC uptake, growth hormones, secretion of digestive enzymes and hormones. The CC sequences of the invention are used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate ZSIG33 expression.

CC The nucleic acids of the invention and their complements are used as DNA CC probes in diagnostic assays to detect and quantitate the presence of Similar nucleic acids in samples, and therefore which patients may be in complements are used as DNA CC probes in the production of antibodies against ZSIG33 apptides are used as antigens in the production of antibodies against ZSIG33 and in assays to identify CC modulators of ZSIG33 expression and activity. The anti-ZSIG33 antibodies are used as diagnostic agents for detecting the presence of ZSIG33 in samples (e.g. by enzyme linked immunosorbent CC anti-ZSIG33 antibodies are also used as diagnostic agents for detecting the presence of ZSIG31 in samples (e.g. by enzyme linked immunosorbent CC assay (ELISA)). The paptides and nucleic acids of the invention are used to modulate gastric contractility, nutrient uptake, growth hormones, the secretion of digestive enzymes and/or secretion of enzymes and/or hormones in the pancreas. zsig33-like DNA is used in gene therapy CC and zsig33-like peptide is used in protein therapy. The present sequence cis human zsig33-like peptide, zsig33-gamma peptide
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growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZSIG33-Like peptides and polynucleotides, useful for modulating gastric contractility, nutrient uptake, growth hormones and/or secretion of digestive/pancreatic enzymes and hormones.
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(SHEP/)
(DEIS/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to zsig33-like peptides and their corresponding
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SHEPPARD P O.
DEISHER T A.
BISHOP P D.
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                   The invention relates to zsig33-like peptides (ZS33LP) including zsig33-CCC linker, zsig33-beta, zsig33-gamma, zsig-33-delta and zsig33-epsilon competides and nucleic acid molecules encoding such zsig33-like peptides. CC ZS33LP peptides activate the immune system in boosting immunity to immunodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines and in communodeficiency virus (HIV) patients, in improving vaccines of the communication of the communication of the communication of the communication enhancers for the computation release. They are also useful as research reagents for the computation, good trigger and or the communication, confirmed of the communication and/or confirmed with cast varieties in all or confirmed with cast varieties.
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Matches 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptides, useful for modulating gastric contractility, nutrient uptake, pancreatic secretion of hormones, digestive enzymes and treatin gastrointestinal and growth related diseases, comprises zsig33-like
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Pred. No. 1.8e+03;
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The invention relates to an isolated pro-apoptotic polypeptide, designated DIABLO, or its biologically active fragment of 8 amino acids in length. Also included are the polypuclectide encoding DIABLO, expression vectors, transformed host cells, producing a biologically expression vectors, transformed host cells, producing a biologically active fragment of DIABLO (by contacting an inhibitor of apoptosis (IAP) with a fragment of the polypeptide, and detecting a reduction in activity of the IAP), producing a natural or synthetic variant of DIABLO with cell
                                                                                                                                                                                                                                                                                                                  New pro-apoptotic polypeptide, useful for screening for agents which modulate cell death and for treating conditions associated with cell death or apoptosis e.g. cancer.
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CC DIABLO in a biological sample (by contacting the sample with an IAP and CC detecting the presence of an IAP/DIABLO complex), modulating the death of CC a cell (by contacting a cell with an agent, which modulates the level CC and/or functional activity of a polypeptide), modulating the death of CC and/or functional activity of a polypeptide), a composition for CC treatment/prophylaxis of a DIABLO related condition comprising an agent which reduces the level/activity of a polypeptide or DIABLO, or a CC nucleic acid encoding DIABLO, is useful for screening for an agent which CC modulates cell death. An antigen-binding molecule is useful for detecting CC DIABLO in a biological sample. The agent which modulates the level and/or functional activity of a polypeptide comprising mature or pro-human CC DIABLO polypeptide, is useful for the treatment and/or prophylaxis of a condition associated with expression or activation of DIABLO, such as CC cancer, vascular disease, hepatic disease, autoimmune disease and CC neurodegenerative disease, hepatic disease, autoimmune disease and CC associated with heart attack, or hepatic tissue damage associated with a CC conditions associated with cell death or apoptosis. The present sequence CC represents a partial peptide sequence from human DIABLO, which co-cc procipitates with the human IAP protein MIHA (not defined)
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                                                                                                                                                                                                                                                                                                                                                                             Gastrointestinal-Gen.; Osteopathic; Anorectic; Metabolic; Endocrine-Gen.; Nephrotropic; Hepatotropic; Respiratory-Gen.; Nootropic; Dermatological; Gynecological; Vulnerary; Antinflammatory; Antinicrobial; Anti-HIV; Antibacterial; Virucide; Cytostatic; Immunomodulatory; Immunostimulant; Gene Therapy; zsig33; musculoskeletal disease; endocrine disease; nutritional disorder; metabolic disorder; growth disorder; wound healing; genitourinary disease; gynecology and obstetrics; renal disease; gynecology and obstetrics; infectious disease;
                                                  N-PSDB;
                                                                                                     Jaspers SR,
                                                                                                                                                                        11-MAY-2000; 2000US-0203300P
10-MAY-2001; 2001US-00853253
                                                                                                                                                                                                                           19-AUG-2004; 2004US-00921371.
                                                                                                                                                                                                                                                               03-MAR-2005
                                                                                                                                                                                                                                                                                                 US2005048618-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human zsig33-gamma peptide, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADY39273 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                       (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                    2005-195284/20
DB; ADY39272.
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3; Conserv
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                                                                                                   Sheppard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                     PO,
                                                                                                         Deisher TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 28; DB
Pred. No. 1.8e
5; Mismatches
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                                                                                                         Bishop
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New zsig33-like polypeptide, useful for treating a disease, e osteoporosis, obesity, dwarfism, wound healing, inflammation,

n, e.g.

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ABF38807
ID ABF3
XX AEF3
XX AEF3
XX AEF3
XX E. C. XX
XX Esch
XX Esch
XX WO20
XX Esch
XX WO20
XX O8-1
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loc
Matches
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This invention describes a novel protein construct comprising a LacI repressor peptide and a His3-20 linker peptide sequence or a LacI repressor peptide (AEF38804), its functional variant and a linker per AEF38810) adapted to bind to support surface. The invention also describes; a) a nucleic acid encoding the construct; b) a vector comprising the nucleic acid and c) a kit comprising the construct in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E. coli LacI repressor hinge region SEQ ID NO
                                                                                                                                                                                                                                                                                       Novel protein comprising LacI repressor peptide and sequence, useful for purifying DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-JUL-2004; 2004GB-00014987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-MAR-2005; 2005WO-GB000888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fusion protein; LacI repressor; reporter protein; DNA purification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-MAR-2006
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                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2006-090395/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                Hine AV, Darby RAJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYAS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEF38807 standard; peptide; 17
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                                                                                                                                                                                                                                SEQ
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                                                                                                                                                                                                                         ID NO 15; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 28; DB 9;
Pred. No. 1.8e+03;
3; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15
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                                                                                                                                                                                                                                                                                                                    a 3-20 linker peptide
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                                                                                                 peptide
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RESULT 14
AAW58715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC lyophilized form and a support surface. The construct comprises a fusion CC protein, in which the LacI repressor peptide is fused, and thus CC functionally bound to the linker peptide. The LacI repressor peptide CC comprises an NH 2 -terminal headpiece (AEF38806), a hinge region CC (AEF38807), an inducer binding domain (AEF38806), and a COOH-terminal CC helix (AEF38809). The construct also comprises a reporter protein CC (AEF38813). The product of the invention is useful for purifying DNA CC comprising at least a region of lacO operator or its variant which CC comprising at least a region of lacO operator sequence CC with the construct to form a DNA/protein complex, attaching the DNA from the CC DNA/protein complex to a support surface, and eluting the DNA from the CC DNA/protein complex. The DNA to be purified comprises a) at least one of CC lacOl, lacOl sites, or its derivative or variant; b) is contained CC in a plasmid or a crude cell lysate. The support surface is adapted to CC bind to the construct and comprises a porous resin. Eluting is carried CC with an allosteric inducer of the LacI repressor protein (e.g. isopropylbeta-D-thiogalactopyranoside or allolactose) in conjunction with salt. CC prior to the elution from the support surface. The construct allows rapid CC and effective isolation of highly pure DNA, and is cost effective. This sequence represents the E. coli LacI repressor protein hinge region, used to make the construct LacI-His6-GFP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                      22-DEC-1989;
09-MAY-1990;
27-AUG-1990;
24-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                 Cytotoxic lymphocyte maturation factor; CLMF; cytokine; synergize; interleukin 2; II-2; human; lymphoblastoid cell; detection; purifying; proliferation; cytotoxic T cell; transplantation; antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW58715;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
                                                            WPI; 1998-413150/35
                                                                                            Podlaski FJ,
Chizzonite RA,
                                                                                                                                                                          02-MAR-1994;
                                                                                                                                                                                                                                                                      02-JUN-1995;
                                                                                                                                                                                                                                                                                                     14-JUL-1998
                                                                                                                                                                                                                                                                                                                                    US5780597-A.
                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tryptic 40 kD subunit CLMF peptide off PVDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW58715 standard; peptide; 17
                                                                                                                                           (HOFF ) HOFFMANN LA ROCHE INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                          Stern AS, C
, Gubler UA;
                                                                                                                                                                          94US-00205011.
                                                                                                                                                                                         90US-00572284.
92US-00857023.
                                                                                                                                                                                                                        90US-00520935
                                                                                                                                                                                                                                       89US-00455708
                                                                                                                                                                                                                                                                      95US-00460061
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                                                                                                           Gately
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Pred. No. 1.8e+03;
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                                                                                                           Μ<u>Κ</u>,
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                                                                                                            Pan
                                                                                                           YE,
                                                                                                            Hulmes
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••
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New antibodies to cytotoxic lymphocyte maturation factor - useful for detecting, purifying, and/or blocking proliferation and activation of cytotoxic T cells, such as in transplantation(s).

Example

3; Col

23; 71pp; English

An isolated antibody has been developed which binds specifically to cytotoxic lymphocyte maturation factor (CLMF) and neutralises CLMF induced proliferation by more than 50% in a CLMF dependent T cell growth assay and/or inhibits binding of at least 60% of the factor to phytohaemagglutinin (PHA) activated peripheral blood lymphocyte (PBL) blasts as determined in a CLMF receptor binding assay. CLMF is a chetrodimeric protein having a molecular weight band of 75 kD, determined by sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) under non-reducing and/or reducing conditions, providing a first subunit having a molecular weight band of 40 kD and a second subunit having a molecular weight of 35 kD. The present sequence represents a tryptic 40 kD CLMF peptide off polyvinylidene difluoride (PVDF), from an example of the present invention. The antibady can be used for the purification chand/or detection of CLMF. It is bedy seed in therapeutic treatments which require selective blocking of proliferation and activation of cytotoxic T cells (CTLs) such as in transplantation

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RESULT 15
AAU78549
ID AAU78549
XX AAU78
AC AAU78
XX Scram
XX Perle
KW Perle
KW Pepax
KW Carti
XX Synth
XX Synth
XX 27-AL
XX 27-AL
XX 27-AL
XX PR 31-AL
PR 17-AL
XX 27-AL
XX 27
                             This invention relates to an adhesive polypeptide derived from domain IV of the extracellular matrix protein perlecan. Perlecan is is a heparin sulphate proteoglycan and is also known as HSPG2. The polypeptide of the
                                                                                                                                                                                                                                                                        New polypeptides derived from the perlecan core protein, useful for strong and selective cell adhesion, particularly as coating for scaffolds for tissue repair, e.g. for treating tooth or cartilage defects, or repairing damaged skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-AUG-2000; 2000US-0229222P.
01-AUG-2001; 2001US-0309209P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             perlecan; cell adhesion; extracellular matrix; adhesive polypeptide; heparin sulphate proteoglycan; HSPG2; scaffold; bone; cartilage; tissue damage; tissue repair; bone loss; tooth loss; gum disease; cartilage defect; rheumatoid arthritis; osteoarthritis; ulcer.
                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-AUG-2001; 2001WO-US026640
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Farach-Carson MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scrambled human perlecan cell adhesion peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002-304245/34.
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                                                                                                                                                                                                Page 18; 41pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carson
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                                                                                                                                                                                                    English.
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                                                                                                                                                                                                                                            invention allows strong and selective cell adhesion and is useful as coating for scaffolds, e.g. implantable tissue scaffolds. The polypeptide may be coated on matrices implanted in damaged tissue (e.g. bone or cartilage) to increase the adhesion of desirable biological materials to the repair site. The scaffolds are useful for treating tooth and jaw defects (e.g. in cases of trauma, bone loss, tooth loss or gum disease) or cartilage defects (e.g. those that result from rheumatoid arthritis, osteoarthritis or trauma), or to repair defects and damage in skin, burns or ulcers. The present sequence represents a peptide composed of the scrambled sequence of human perlecan adhesion peptide. This peptide was used as a control for cell adhesion assays in the examples of the method of the inverser.
Sequence 17 AA;
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유 8 Best Matches Local w 4 RSMGGKEDLIW 14 Similarity 6; Conserv :|:|| ||| KSVGG--SLIW 11 Conservative 31.4%; 2 Score 27; Pred. No. Mismatches 2.6e+03; 'n Gaps

Query Match

DB 5;

Length 17

Search co Job time completed: June 13, 198 secs 2006, 01:27:39

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Query Match
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31.4%;

Score 27; DB 2; Pred. No. 2.6e+03;

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Mismatches

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Sequence 17 AA;

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cytochrome c554 -	T cell receptor be	T cell receptor al	Ig heavy chain DJ	Ig heavy chain DJ	Ig heavy chain DJ	Ig heavy chain CRD	C	hypothetical prote	plastocyanin - Ara	heterogeneous ribo	enamelin, 26K - bo	dihydrolipoamide S	unidentified prote	Ig H chain V-D-J r	ribosomal protein

C;Superfamily: histone H4 C;Keywords: acetylated amino end; acetyllysine F;1/Modified site: acetylated amino end (Ser) #status experimental F;5,12,16/Binding site: acetyl (Lys) (covalent) (partial) #status experimental A;Status: preliminary A;Molecule type: protein A;Residues: 1-17 <WOU> C;Accession: A61321 R;Wouters-Tyrou, D.; Martin-Ponthieu, A.; Sautiere, P.; Biserte, G. FEBS Lett. 128, 195-200, 1981 A;Title: Acetylation of histone H4 in chicken erythrocyte and cuttle-fish testis chromat A;Reference number: A61321; MUID:81261219; PMID:7262310 A;Accession: A61321 histone H4 - common cuttlefish (fragment) C;Species: Sepia officinalis (common cuttlefish) C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004 RESULT A61321 A;Molecule type: protein A;Residues: 1-17 <EST'> A;Residues: 1-17 <EST'> A;Cross-references: UNIPROT:07M432; UNIPARC:UPI0000175C3B C;Superfamily: trypsin; trypsin homology C;Keywords: hydrolase; protein digestion; serine proteinase R;Estell, D.A.; Laskowski Jr., M. Biochemistry 19, 124-131, 1980 A;Title: Dermasterias imbricata trypsin 1: an enzyme which rapidly hydrolyzes the reacti A;Reference number: A61334; MUID:80109692; PMID:7352972 A;Accession: B61334 trypsin (EC 3.4.21.4) 2 - starfish (Dermasterias imbricata) (fragment) C;Species: Dermasterias imbricata C;Species: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul A; Cross-references: UNIPROT: Q7M3Z5; UNIPARC: UPI00001771C6 RESULT Accession: B61334 Matches Query Match Best Local Query Match Best Local Similarity 6 MGGKE 10 Similarity 4. 4. VGGKE 6 Conservative Conservative 24.48; 26.7%; ٥, Score 23; DB 2; Pred. No. 1.6e+03; 1; Mismatches (Score 21; DB 2; Pred. No. 3.5e+03; 0; Mismatches 2 17-Jul-1994 #text_change 09-Jul-2004 . . Length 17 Length 17; Indels 0 0 Gaps Gaps 0 0

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hypothetical protein Z4331 [imported] - Escherichia coli (strain O157:H7, substrain C;Species: Escherichia coli C;Species: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004 C;Accession: G85956
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-17 <STO>
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G85956
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C;Species: 17-Uul-1994 #sequence_revision 17-Jul-1994 #text_change 31-Dec-2004
C;Accession: B61321
R;Wouters-Tyrou, D; Martin-Ponthieu, A.; Sautiere, P.; Biserte, G.
FEBS Lett. 128, 195-200, 1981
A;Title: Acetylation of histone H4 in chicken erythrocyte and cuttle-fish testis chromat A;Reference number: A61321; MUID:81261219; PMID:7262310
A;Accession: B61321
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-17 < MUID:81261219; PMID:7262310
                                                                  A; Accession: G85956
                                                                                     A;Title: Genome sequence of enterohemorrhagic Escherichia coli
A;Reference number: A85480; MUID:21074935; PMID:11206551
                                                                                                                                  C;Accession: G85956
R;Perna, N.T.; Plunkett III, G.; l
iller, L.; Grotbeck, E.J.; Davis,
Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Experimental source: T lymphocyte C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R.Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L. J. Exp. Med. 174, 1371-1383, 1991
A.Title: T cell receptor genes in a series of class I major histocompatibility complex-sallelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T-cell receptor alpha chain (RA10.3.3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0778
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C;Keywords: acetylated amino end; acetyllysine
F;1/Modified site: acetylated amino end (Ser) #status experimental
F;5,12,16/Binding site: acetyl (Lys) (covalent) (partial) #status:
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protein Pfd3 - golden needle mushroom (fragment)
C;Species: Flammulina velutipes (golden needle mushroom)
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text
C;Accession: E59137
                                                                                       RESULT
E59137
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A;Status: preliminary; not compared with conceptual translation A;Molecule type: nucleic acid A;Residues: 1-17 <ROS>
                                                                                                                                                                                                                                                               R;Rosenberg, W.M.; Moss, P.A.; Bell, J.I.
Eur. J. Immunol. 22, 541-549, 1992
A;Title: Variation in human T cell receptor V beta and J beta repertoire: analysis using A;Reference number: A49039; MUID:92164737; PMID:1311263
                                                                                                                                                                                                                                                                                                                                      T-cell receptor beta chain V-D-J-C region (V beta 8, J beta 2.7) - human (f C;Species: Homo sapiens (man) C;Cpecies: Homo sapiens (man) C;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 30-May-1997 C;Accession: B49255
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A;Molecule type: mRNA
A;Residues: 1-17 <POR>
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R;Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, S.P.
J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR) expression
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PH1789
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A;Experimental source: strain O157:H7, substrain EDL933
C;Genetics:
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                                                                                                                                        A;Note: sequence extracted from C;Keywords: T-cell receptor
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A;Note: sequence extracted from NCBI backbone (NCBIP:90723)
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C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change
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7.7e+03;
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5.2e+03;
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#text_change

09-Jul-2004

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us-10-612-162a-1.exact.rpr
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W;Alternate names: conotoxin GV; sleeper peptide
C;Species: Conus geographus (geography cone)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: A05168
R;McIntosh, J.M.; Olivera, B.M.; Cruz, L.J.; Gray, W.R.
J. Biol. Chem. 259, 1433-14346, 1984
A;Title: gamma-Carboxyglutamate in a neuroactive toxin.
A;Reference number: A05168; MUID:85054897; PMID:6501296
A;Accession: A05168
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A;Description: Differences of proteins expressed in the fruiting dikaryon and the non-fu A;Reference number: A59137
A;Recession: E59137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Accession: A61334
R;Estell, D.A.; Laskowski Jr., M.
R;Estell, D.A.; Laskowski Jr., M.
Biochemistry 19, 124-131, 1980
A;Title: Dermasterias imbricata trypsin 1: an enzyme which rapidly hydrolyzes
A;Reference number: A61334; MUID:80109692; PMID:7352972
A;Reference number: A61334; MUID:80109692; PMID:7352972
                                                                                      A;Accession: A61334
A;Molecule type: protein
A;Residues: 1-17 <EST>
                                                                                                                                                                                                                                                          trypsin (EC 3.4.21.4) 1 - starfish (Dermasterias imbricata) (fragment)
C;Species: Dermasterias imbricata
C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Keywords: amidated carboxyl end; carboxyglutamic acid; neurotoxin; venom F;3,4,7,10,14/Modified site: gamma-carboxyglutamic acid (Glu) #status experimental F;17/Modified site: amidated carboxyl end (Asn) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Contents: annotation; conformation by (1) H-NMR C; Superfamily: conantokin
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C;Superfamily: trypsin; trypsin homology
                                             A;Cross-references: UNIPROT:Q7M433; UNIPARC:UPI0000175C3A
C;Comment: This isoform is novel in that it readily dissociates from bovine
                                                                                                                                                                                                                                                                                                                                   A61334
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Best Local
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2; Conserv
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Pred. No. 7.7e
3; Mismatches
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7.7e+03;
2;
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pollen allergen Fes e I type A - reed fescue (fragment) C;Species: Festuca elatior (reed fescue) C;Date: 28-Apr-1993 #sequence_revision 28-Apr-1993 #tex C;Accession: C37396 R;Esch, R.E.; Klapper, D.G. Mol. Immunol. 26, 557-561, 1989
                                                                                                          RESULT 13
C37396
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A61019
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A; Note: sequence extracted from NCBI backbone
C; Superfamily: caldesmon
                                                                                                                                                                                                                                                                                                                                                                                                Histochemistry 93, 513-518, 1990
A;Title: N-terminal amino acid sequence, immunohistochemical A;Reference number: A61019; MUID:90236771; PMID:2332352
A;Accession: A61019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Rattus norvegicus (Norway rat)
C;Date: 31-Dec-1993 #sequence_revision 31
C;Accession: A61019
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A;Title: Determination of the phosphorylation sites of smooth muscle A;Reference number: A44873; MUID:91378498; PMID:1898046
A;Accession: B44873
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A;Molecule type: protein
A;Residues: 1-17 <IKE>
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R;Ikebe, M.; Hornick, T.
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    Mismatches

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Pred. No. 1.1e+04;
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Pred. No. 1.1e+04;
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Pred. No. 1.1e+04;
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C;Species: Homo sapiens (man)
C;Date: 01-Nov-1996 #sequence_revision 01-Nov-1996 #text_change 05-Nov-1999
C;Accession: A49635
R;Pribyl, T.M.; Campagnoni, C.W.; Kampf, K.; Kashima, T.; Handley, V.W.; McMahon, J.; Caproc. Natl. Acad. Sci. U.S.A. 90, 10695-10699, 1993
Proc. Natl. Acad. Sci. U.S.A. 90, 10695-10699, 1993
A;Title: The human myelin basic protein gene is included within a 179-kilobase transcrip A;Reference number: A49635; MUID:94068468; PMID:7504278
A;Accession: A49635
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C30221

histone H2A.8 - chicken (fragment)
c;Species: Gallus gallus (chicken)
c;Species: Gallus gallus (chicken)
c;Apate: 02-Nov-1989 #sequence_revision 02-Nov-1989 #text_change 30-Sep-1993
C;Accession: C30221
R;Challoner, P.B.; Moss, S.B.; Groudine, M.
Mol. Cell. Biol. 9, 902-913, 1989
A;Title: Expression of replication-dependent histone genes in avian spermatids involves A;Reference number: A30221; MUID:89261754; PMID:2471062
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-17 <CHA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Isolation and characterization of a major cross-reactive grass group I allerger A;Reference number: A37396; MUID:89364850; PMID:2475768
A;Accession: C37396
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-17 <ESC>
A;Cross-references: UNIPARC:UPI000017B0DA
C;Keywords: pollen
Search completed: June 13, 2006, 01:44:16 Job time : 39 secs
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A;Molecule type: DNA
A;Residues: 1-17 <RES>
A;Cross-references: UNIPARC:UPI000016AFAB; GB:L18862; NID:g435054; PIDN:AAA72008.1;
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A49635
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15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 6.
Trypsin (EC 3.4.21.4) 2 (Fragment).
Dermasterias imbricata (Sea star).
Eukaryota, Metazoa; Echinodermata; Eleutherozo Asteroidea; Valvatacea; Walvatida; Asterinidae NCBI TaxID-7590;
                                PROTEIN SEQUENCE. MEDLINE=96250169;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Dermasterias imbricata trypsin 1: an enzyme which the reactive-site peptide bonds of protein trypsin Biochemistry 19:124-131(1980).
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71.4%;
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Pred. No. 1.
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Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4C7B64410E84D968 CRC64;
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Asterinidae; Dermasterias
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1.9e+04;
hes 0; Indels
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inhibitors.";
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RESULT 6
Q85AH6 9PRIM
ID Q85AH6;
AC Q85AH6;
DT 01-JUN-2003, in
DT 07-FEB-2006, en
DT 07-FEB-2006, en
DE Cytochrome oxid
GN Name=COXII;
OS Lepilemur edwar
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SEQUENCE
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P82697;
04-MAY-2001, integrated into UniProtKB/Swiss-Prot
01-MAX-2001, sequence version 1.
07-FEB-2006, entry version 21.
Peptide hormone 4 (Pea-YLS-amide).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ferrer P., Hedegaard L., Halkier T., Diers I., Savva D "Molecular cloning of a lytic beta-1,3-glucanase gene xanthineolytica LLG109. A beta-1,3-glucanase able to s permeabilize the yeast cell wall.";
Ann. N. Y. Acad. Sci. 782:555-565(1996).
                            01-JUN-2003, integrated into UniProtKB/TrEMBL.
01-JUN-2003, sequence version 1.
07-FEB-2006, entry version 12.
Cytochrome oxidase subunit III (Fragment).
                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/cerms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Periplaneta americana (American cockroach).
Bukaryota; Metazoa; Arthropoda; Haxapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; E
Blattidae; Blattinae; Periplaneta.
               Cytochrome oxidase Name=COIII;
                                                                                                                                                                                                                                                                                                                                    PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                               of insects.";
Ann. N. Y. Acad. Sci. 897:282-290(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROTEIN SEQUENCE, AND MASS SPECTROMETRY.
TISSUE-Abdominal perisympathetic organs;
MEDLINE=20140865; PubMed=10676456;
Predel R., Eckert M., Holman G.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                   Amidation;
   Lepilemur edwardsi (Milne-Edwards's sportive lemur).
                                                                                                                                                                                                                                                                                                                                                                                                                          -!- MASS SPECTROMETRY: MW=1966.48; METHOD=MALDI; RANGE=1-17;
NOTE=Ref.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The unique neuropeptide pattern in abdominal perisympathetic organs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=6978;
                                                                                                                                                                                                                                               Local
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3; Conserv
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1 17 Peptide hormone 4.
PFIde-PRO 0000044205.
17 17 Serine amIde.
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                                                                                                      PRELIMINARY;
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75.0%;
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Pred. No. 1.9e
1; Mismatches
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                                                                                                                                                                                                                                   2; Mismatches
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Pred. No.
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se gene from Oerskovia
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Blattoidea;
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Best Local S
Matches 3
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Q85AH7;
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NON TER 1
SEQUENCE 17 A
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EMBL;
GO; GC
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Lemuridae; Hapalemur.
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90; 99
90; 99
90; 99
         EMBL; AF224575; AAO37906.1; -; Genomic_DNA.
EMBL; AF224576; AAO37910.1; -; Genomic_DNA.
EMBL; AF224577; AAO37914.1; -; Genomic_DNA.
                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                      Pastorini J., Forstner M.R.J., Martin "Phylogenetic relationships of gentle Evol. Anthropol. 11:150-154(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL;
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Mammalia; Eutheria; Euarc
Megaladapidae; Lepilemur.
MCBI_TaxID=122230;
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                                                                                                                                                                                                                                                                                    Hapalemur griseus alaotrensis
                                                                                                                                                                                                                                                                                                Name=COIII;
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                                                                                                   "A molecular approach lemurs.";
                                                                                                                         Pastorini J.,
                                                                                                                                    MEDLINE=22631663;
                                                                                                                                               NUCLEOTIDE SEQUENCE
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L; AF24594; AAF33640.1; -; Genomic DNA.
GO:0016020; C:membrane; IEA.
GO:0005739; C:mitochondrion; IEA.
GO:0004129; F:cytochrome-c oxidase activity;
GO:0006118; F:electron transport; IEA.
erPro; IPR000298; CytC oxdse III.
THER; PTHR11403; CytC oxdse III; 1.
m. PEF01510, COY3.
 GO:0016020;
                                                                                       Natl. Acad. Sci. U.S.A. 100:5879-5884(2003)
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                                                                                                                          Thalmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    663; PubMed=12719521; DOI=10.1073/pnas.1031673100; Thalmann U., Martin R.D.;
                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
 C:membrane;
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                                                                                                                                   PubMed=12719521; DOI=10.1073/pnas.1031673100;
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2132 MW;
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Euarchontoglires; Primates; Strepsirhini;
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III
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Pred. No.
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lemurs
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                                                                                                                                                                                (Hapalemur).";
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Q85DA6 GALCR Q85DA6 GALCR ID Q85DA6; AC Q85DA6; DT 01-JUN-2003, DT 01-FEB-2006;

entry version 11

integrated into UniProtKB/TrEMBL

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A2 GALSE
Q85DA2 GALSE
Q85DA2;
                                                                                                                                      EMBL; AF224644; AAP33660.1; -; Genomic_DNA. GC; GO:0016020; C:membrane; IEA. GO; GO:0005739; C:mitochondrion; IEA. GO; GO:0004129; F:cytochrome-c oxidase acti: GO; GO:0004129; F:electron transport; IEA. GO; GO:0006118; P:electron transport; IEA. InterPro; IPR00298; CytC_oxdse_III. PANTHER; PTHR11403; CytC_oxdse_III; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GG; GO:0005739; C:mitochondrion; IEA.
GG; GO:0004129; F:cytochrome-c oxidate
GG; GO:0006118; P:electron transport; I
InterPro; IPR000298; CytC_oxdse_III.
PANTHER; PTHR11403; CytC_oxdse_III; 1.
                                                                                                                                                                                                                                                                                                   MEDLINE=22631663; PubMed=12719521; DOI=10.1073/pnas.1031673100; Pastorini J., Thalmann U., Martin R.D.; "A molecular approach to comparative phylogeography of extant M
                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
Galagonidae; Galago.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-FEB-2006, entry version 11. Cytochrome oxidase subunit III
                                                                                             SEQUENCE
                                                                                                                    Mitochondrion.
                                                                                                                                Pfam; PF00510; COX3;
                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
MEDLINE=22631663; Pu
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                                              Conservative
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                                                         25.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Mismatches
                                             Score 22; DB 2; Le
Pred. No. 2.8e+04;
2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 22; DB 2;
Pred. No. 2.8e+04;
                                                                                             CBEB69E354D98978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CBEB69E354D98978 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                              lesser bushbaby).
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RESULT 10
Q85DB4 LEPRU
ID Q85DB4 LEPRU
AC Q85DB4 |
DT 01-JUN-2003, s
DT 07-FEB-2006, e
DE Cytochrome oxid
GN Name=COIII ;
OS Lepilemur rufi
OG Mitochondrion.
OC Eukaryota, Met
OC Eukaryota, Met
OC Megaladapidae;
OX NCBI_TaxID=788
RN NICLEOTIDE SEC
RX MEDLINE=226316
RA PASTOYINI J,
RT Proc. Natl Ac
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Best Local Similarity
Matches 3; Conserv
                                            EMBL; AR234596; AAP33648.1; -; Genomic_DNA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0004129; F:cytochrome-c oxidase activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006118; P:electron transport; IEA.
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Mammalia; Eutheria;
Megaladapidae; Lepil
NCBI_TaxID=78866;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=22631663; PubMed=12719521; DOI=10.1073/pnas.1031673100;
Pastorini J., Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant Malagasy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytochrome oxidase subunit III Name=COIII;
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GO; GO:0016020; C:membrane; IEA.
GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0004129; F:cytochrome-c oxidase activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR000298; CytC_oxdse_III.
PANTHER; PTHR11403; CytC_oxdse_III; 1.
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Pastorini J., Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant M
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                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mitochondrion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proc. Natl. Acad. Sci. U.S.A. 100:5879-5884(2003).
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Pred. No. 2.8e+04;
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                                                                                                       Query Match
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EMBL; AF224586; AAN64793.1; -; Genomic_DNA.
EMBL; AF224587; AAN64797.1; -; Genomic_DNA.
EMBL; AF224587; AAN64797.1; -; Genomic_DNA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0004129; F:cytochrome-c oxidase activi
GO; GO:0006118; P:electron transport; IEA.
Interpro; IPR000298; CytC_oxdse_III.
PANTHER; PTHR11403; CytC_oxdse_III; 1.
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SEQUENCE 17 1
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Mammalia; Eutheria;
Lemuridae; Varecia.
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                                                                                                                                                             SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pastorini J., Forstner M.R.J., Mari
"Phylogenetic relationships of geni
Evol. Anthropol. 11:150-154(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=22631663; PubMed=12719521; DOI=10.1073/pnas.1031673100, Pastorini J., Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOFIDE SEQUENCE.
MEDLINE=22281630; PubMed=12393004; DOI=10.1016/S0047-2484(02)90587-7;
Pastcorini J., Forstner M.R., Martin R.D.;
"Phylogenetic relationships among Lemuridae (Primates): evidence from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q<del>В</del>НВ37_VARVV
Q8НВ37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=COIII;
                                                                                                                                                                                   NON TER
                                                                                                                                                                                                          Mitochondrion.
                                                                                                                                                                                                                              Pfam; PF00510; COX3;
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                                                                                 Local
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                                                          l Similarity
3; Conserv
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            DLIMELL
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                                                       25.6%;
nilarity 42.9%;
Conservative
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                                                                                                                                                          2132 MW;
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                                                          Score 22; DB 2;
Pred. No. 2.8e+04;
2; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Martin R.D.;
gentle lemurs
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                                                                                                            Length 17;
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DVVWLFL

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RESULT 12

QBHB38 VARVR
QBHB38 VARVR
QBHB38,
AC QBHB38,
DT 01-MAR-2
DT 01-MAR-2
DT 07-FEB-2
DT 07-FEB-
RESULT 13

CORHSJ 9 HAPSI PRELIMINARY; PRT; 17 A

AC Q8HB39;
DT 01-MAR-2003, integrated into UniProtKB/TrE

DT 01-MAR-2003, sequence version 1.

DT 07-FEB-2006, entry version 12.

DE Cytochrome oxidase subunit III (Fragment).
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Best Local S
Matches 3
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Distributed
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MEDLINE=22631663; PubMed=12719521; DOI=10.1073/pnas.1031673100;
Pastorini J., Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pastorini J., Forstner M.R.J., Martin "Phylogenetic relationships of gentle Evol. Anthropol. 11:150-154(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
MEDLINE=22281620; PubMed=12393004; DOI=10.1016/S0047-2484(02)90587-7;
Pastcorini J., Forstner M.R., Martin R.D.;
"Phylogenetic relationships among Lemuridae (Primates): evidence from mtDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-2003,
07-FEB-2006,
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QBHB38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Proc. Natl. Acad. Sci. U.S.A. 100:5879-5884(2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
Lemuridae; Varecia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Varecia variegata rubra
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MBL; AF224589; AAN64805.1; -; Genomic_DNA.
MBL; AF224591; AAN64813.1; -; Genomic_DNA.
MBL; AF224590; AAN64810.1; -; Genomic_DNA.
MBL; AF224592; AAN64817.1; -; Genomic_DNA.
MBL; AF224592; AAN64817.1; -; Genomic_DNA.
O; GO:0016020; C:membrane; IEA.
O; GO:0016020; C:membrane; IEA.
O; GO:0004129; F:cytochrome-c oxidase_activi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GO:0006118; P:electron transport; IEA. erPro; IPR000298; CytC_oxdse_III. THER; PTHR11403; CytC_oxdse_III; 1.
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                     <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                   Score 22; DB Pred. No. 2.8e 2; Mismatches
                                                                                UniProtKB/TrEMBL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CBEB69E354D98978 CRC64;
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Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
2.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          activity; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Length 17;
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Q8HB40_HAPGR
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MEDLINE=22281620; Pubmed=12393004; DOI=10.1016/S0047-2484(02)90587-7;
Pastorini J., Forstner M.R., Martin R.D.;
"Phylogenetic relationships among Lemuridae (Primates): evidence from mtDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF224583; AAN64781.1; -; Genomic_DNA.
EMBL; AF224584; AAN64785.1; -; Genomic_DNA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0006129; F:cytochrome-c oxidase activación GO:0006118; P:electron transport; IEA.
InterPro; IPR000298; CytC_oxdse_III.
PANTHER; PTHR11403; CytC_oxdse_III; 1.
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Q8HB40;
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                                                                                                                                  Lemuridae; Hapalemur.
NCBI_TaxID=122221;
                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Primates; Stre
                                                                                                                                                                                                                                      Mitochondrion.
                                                                                                                                                                                                                                                                                 Name=COIII;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
MEDLINE=22281620; PubMed=12393004; DOI=10.1016/S0047-2484(02)90587-7;
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lmann U., Martin R.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2132 MW;
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SEQUENCE

Hum. Evol. 43:463-478(2002).

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RESULT 15
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Q8H941;
Q9H941;
Q1-MAR-2003, integrated into UniProtKB/TrEMBL.
Q1-MAR-2003, sequence version 1.
Q7-FEB-2006, entry version 12.
Q7-FEB-2006 anti-age anti-age of the control of the
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EMBL; AF224579; AAN6479.1; -; Genomic_DNA.
EMBL; AF224580; AAO37918.1; -; Genomic_DNA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005739; C:mitochondrion; IEA.
                                                                                     NUCLEOTIDE SEQUENCE.
MEDLINE=22631663; PubMed=12719521; DOI=10.1073/pnas.1031673100;
Pastcorini J. Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant Malagasy
                                                                                                                                                                                                                                                                                                                                                                 Pastorini J., Forstner M.R.J., Martin "Phylogenetic relationships of gentle Evol. Anthropol. 11:150-154(2002).
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MEDLINE=22281620; PubMed=12393004; DOI=10.1016/S0047-2484(02)90587-7;
Pastorini J., Forstner M.R., Martin R.D.;
"Phylogenetic relationships among Lemuridae (Primates): evidence from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Strepsirhini;
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GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0004129; F:cytochrome-c oxidase activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR000298; CytC_oxdse_III.
PANTHER; PTHR11403; CytC_oxdse_III; 1.
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Pastorini J., Thalmann U., Martin R.D.;
"A molecular approach to comparative phylogeography of extant Malagasy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hapalemur aureus (Golden bamboo lemur).
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GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006118; CytC oxdse_III.
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PANTHER; PTHR11403; CytC_oxdse_III; 1.
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AF224582; AAN64777.1; -; Genomic_DNA.
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3; Conserv
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                                                  2006, 01:32:34
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Pred. No.
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Searched:
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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23.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Issued Patents AA:*

1. /EMC_Celerra_SIDS3/ptodata/2/iaa/5_COMB.pep:*

2. /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

3. /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*

4. /EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*

5. /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*

6. /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7. /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
 DB
                                                   US-09-853-253-14
US-08-497-599-5
US-09-664-677A-18
US-09-664-677A-18
US-09-695-437A-20
US-09-695-769B-229
US-09-490-153-229
US-09-490-324-229
US-09-490-324-229
US-08-635-928-33
US-08-635-928-33
US-08-635-928-33
US-08-382-013A-9
US-08-382-013A-9
US-08-381-625-52
US-08-381-625-52
US-08-381-625-52
US-08-497-599-4
US-08-181-083-52-1
US-09-181-083-52-1
US-09-181-083-52-1
US-09-635-705-21
US-09-635-705-21
US-09-635-705-21
US-09-635-705-21
US-09-635-705-21
US-09-750-754-52
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(without alignments)
31.000 Million cell updates/sec
             Sequence 14, Appli
Sequence 18, Appli
Sequence 10, Appli
Sequence 20, Appli
Sequence 229, App
Sequence 229, App
Sequence 229, App
Sequence 230, Appli
Sequence 33, Appli
Sequence 14, Appli
Sequence 16, Appli
Sequence 21, Appli
Sequence 22, Appli
Sequence 23, Appli
Sequence 23, Appli
Sequence 24, Appli
Sequence 25, Appli
Sequence 27, Appli
Sequence 21, Appli
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Sequence
58,
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US-08-569-188-14	US-08-569-188-4	US-08-105-416-29	US-08-076-092-10	5266328-7	PCT-US94-05684-14	PCT-US94-02631-24	PCT-US91-05177-2	US-09-428-082B-204	US-08-753-750B-55	US-08-751-305-18	US-08-497-599-13	US-08-485-508-116	US-08-741-599-3	US-08-438-883-3	US-08-704-170-24	US-08-438-759-14	US-08-370-567-14	US-08-241-054-116
Sequence 14, Appl	Sequence 4, Appli	Sequence 29, Appl	Sequence 10, Appl	Patent No. 5266328	Sequence 14, Appl	Sequence 24, Appl	Sequence 2, Appli	Sequence 204, App		Sequence 18, Appl	Sequence 13, Appl	Sequence 116, App	Sequence 3, Appli	Sequence 3, Appli	Sequence 24, Appl	Sequence 14, Appl	Sequence 14, Appl	Sequence 116, App

RESULT 1 US-09-853-253-14 US-09-853-253-14 Sequence 14, Application US/09853253 Patent No. 6897286 GENERAL INFORMATION: APPLICANT: JASPERS, STEPHEN APPLICANT: BISHOR, PAUL APPLICANT: BISHOR, PAUL APPLICANT: BISHOR, PAUL TITLE OF INVENTION: Zsig33-like Peptides FILE REFERENCE: 00-30 CURRENT APPLICATION NUMBER: US/09/853,253 CURRENT FILING DATE: 2001-05-10 PRIOR APPLICATION NUMBER: 60/203,300 PRIOR FILING DATE: 2000-05-11 NUMBER OF SEQ ID NOS: 28 SOFTWARE: FASTSEQ for Windows Version 3.0 SEQ ID NO 14 LENGTH: 17 TYPE: PRT ORGANISM: Homo sapiens

Query Match Best Local Similarity 50.0%; Pred. No. 3.4e+02; Matches 3; Conservative 3; Mismatches 0; Indels 0; Gaps Qy
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TELEFAX: (202) 408-4400
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 17 amino acids
TYPE: amino acid
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-497-599-5
 문
                                                                                                                                             ; ORGANISM: Human Immunodeficiency Virus (HIV-1) US-09-564-677A-18
                                                                                                                                                                                                                                                                                                                          APPLICANT: Assaf Friedler
APPLICANT: Abraham Loyter
APPLICANT: Abraham Loyter
APPLICANT: Chaim Gilon
APPLICANT: Amnon Wolf
TITLE OF INVENTION: Inhibition of Nuclear Imp
FILE REFERENCE: 2254-019-99
CURRENT APPLICATION NUMBER: US/09/564,677A
CURRENT FILING DATE: 2000-05-04
PRIOR APPLICATION NUMBER: 122343
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                                                                                                                                                                                        PRIOR FILING DATE: 1997-11-27
PRIOR APPLICATION NUMBER: PCT/IL98/00577
PRIOR FILING DATE: 1998-11-26
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18
LENGTH: 17
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Best Local Similarity 28.6
                                                                                       Query Match
Best Local :
                                                                        Matches
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Patent No. 6664368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: TOOHEY, KIMDELLL.
REGISTRATION NUMBER: 35,391
REFERENCE/DOCKET NUMBER: 02481.1451-00000
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/497,599
FILING DATE: 30-JUN-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                      Similarity 5; Conserv
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Pred. No. 7.5e+02;
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RESULT 4

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CURRENT APPLICATION NUMBER: US/09/845,160
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: JP 2001-131688
PRIOR FILING DATE: 2001-04-27
PRIOR APPLICATION NUMBER: JP 2000-161577
PRIOR FILING DATE: 2000-05-31
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 4
LENGTH: 17
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Best Local Similarity
Matches 3; Conserve
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US-09-845-160-4
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                                                                                                             US-09-695-437A-40
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                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1993-10-06
NUMBER OF SEQ ID NOS: 64
SOFTWARE: PatentIn version 3.1
SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 40, Application US/09695437A Patent No. 6803203 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/09845160 Patent No. 6921663
                                    Matches
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                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: BSA 01-02
CURRENT APPLICATION NUMBER: US/09/695,437A
CURRENT FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 08/398,139
PRIOR FILING DATE: 1995-03-03
PRIOR APPLICATION NUMBER: 08/132,284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Brookhaven Science Associates
APPLICANT: Anderson, Carl W
APPLICANT: Connelly, Margery A
TITLE OF INVENTION: DNA-PK Assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: MIZUGUCHI, HIROYUKI
APPLICANT: HAYAKAWA, TAKAO
TITLE OF INVENTION: ADENOVIRUS VECTOR
                                                                                                                      NAME/KEY: MISC_FEATURE
LOCATION: (1)..(4)
OTHER INFORMATION: human hsp90 residues 1-4
NAME/KEY: MISC_FEATURE
LOCATION: (5)..(17)
OTHER INFORMATION: human p53 residues 15-27 with S20E substitution
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                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                         FEATURE:
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                                                         Local
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6; Conserv
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                                      Conservative
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75.0%;
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                                                       Score 24.5;
Pred. No. 1
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Pred. No. 1.1e+03;
                                      Mismatches
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                                                       5; DB 2;
1.4e+03;
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                                                                      Length 17;
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RESULT 7
US-09-490-070A-229
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US-09-025-769B-229
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                                                                                                                                                                                  Patent No. 6696248
GENERAL INFORMATION:
                                                                                                                                                                                                                         Sequence 229, Application US/09490070A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 229, Application US/09025769B Patent No. 6300064
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Knappi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-AUG-1995 ATTORNEY/AGENT INFORMATION: NAME: James F. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794 REFERENCE/DOCKET NUMBER: MORPHO/5 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
Admes F. Haley, Jr., Esq. c/o Fish & Neave STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ge, Liming
APPLICANT: Moroney, Simon
APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                       y Match 27.9%;
Local Similarity 43.8%;
hes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                Moroney, Šimon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                  APPLICANT: Knappik, Achim
                                                                                                                                                                                                                                                                                                                                                      3 ARSM----GGKEDLIW 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 amino acids
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Ilag, Vic
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                                                                                                                            Ilag, Vic
                                                                                                                                             Pack, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        internal
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Colin G. Sandercock, Esq. c/o Heller Ehrman
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Pred. No. 1.7e+03;
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US-09-490-153-229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 229, Application Patent No. 6706484
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 912-2020
INFORMATION FOR SEQ ID NO: 229:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Colin G. Sandercock, Esq.
REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 37629-0005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 912-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               y Match 27.9%;
Local Similarity 43.8%;
hes 7; Conservative
COUNTRY: USA

ZIP: 10021

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/490,153

FILING DATE: 24-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/490,070A
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. Haley,
STREET: 1251 Avenue of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein FRAGMENT TYPE: internal SEQUENCE DESCRIPTION: SEQ ID NO: 229:
                                                                                                                                                                                                                                                                                                                    Moroney, Simon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Knappik, Achim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin. Release #1.0, Version #1.30 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1666 K Street, N.W., Suite 300 CITY: Washington
                                                                                                                                                                                                                STATE: New York
                                                                                                                                                                                                                                      CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
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Pred. No. 1.7e+03;
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Americas
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ATTORNEY/AGENT INFORMATION:

APPLICATION NUMBER: US/09/025,769B FILING DATE: 18-FEB-1998 APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-AUG-1995

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RESULT 9
US-09-490-324-229
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS
                                                  APPLICATION NUMBER: US/09/490,324
FILING DATE: 24-Jan-2000
PRIOR APPLICATION NUMBER: US/09/025,769
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-FEB-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REFERENCE, DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: James F. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Moroney, Simon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein FRAGMENT TYPE: internal
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STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OF SEQUENCES: 373
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nilarity 43.8%;
Conservative
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                      (212)596-9090
Q ID NO: 229:
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Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                      CLASSIPICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/01302
FILING DATE: 10/27/94
APPLICATION NUMBER: P4337599.5
FILING DATE: 11/01/93
ATTORNEY/AGENT INFORMATION:
NAME: MOLANO, Michael A.
REGISTRATION NUMBER: 39,777
REFERENCE/DOCKET NUMBER: WELT2010(WO/US)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
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APPLICANT: CONRAD
APPLICANT: DINKEL
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MEDIUM TYPE: Diskette
COMPUTER: ISM PC COMPATIBLE
OPERATING SYSTEM: MS-DOS
SOFTMARE: Wordberfect 7.0
CURRENT APPLICATION DATA:
                                                                                                                                                 TELEFAX: 415-362-2928 INFORMATION FOR SEQ ID NO:
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                             SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
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ADDRESSEE: FLIESLER, DUBB, MEYER & LOVEJOY LLP
STREET: Four Embarcadero Center, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
STRANDEDNESS:
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FILING DATE: 09/20/96
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                                                                                                                                                                                                                    TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 ARSM----GGKEDLIW 14
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TYPE: amino acid
STRANDEDNESS: <Unknown>
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SCHLER, HANS MARTIN
SCHER, HANS MARTIN
SCHWEIDER, HANS MARTIN
SCHWEIDER, METAL-BINDING CYSTEIN-FREE
INVENTION: METAL-BINDING CYSTEIN-FREE
INVENTION: METAL-BINDING CYSTEIN-FREE
INVENTION: METHODS FOR THEIR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC Compatible
                                                                                                                                                                                                                       415-362-3800
   single
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                                                                                                                                                    32:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                  TELEFAX: 415-362-2928
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                FILING DATE: 11/01/93
ATTORNEY/AGENT INFORMATION:
ANAME: MOLANO, Michael A.
REGISTRATION NUMBER: 39,777
REFERENCE/DOCKET NUMBER: WBLT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-362-3800
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/
FILING DATE: 09/20/96
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT
FILING DATE: 10/27/94
APPLICATION NUMBER: P43
FILING DATE: 11/01/93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLIESLER, DUBB, MEYER & LOVEJOY LLP
STREET: Four Embarcadero Center, Suite 400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: MALIN, REINHARD
APPLICANT: SCHIER, HANS MARTIN
APPLICANT: SCHNEIDER-MERGENER, JENS
APPLICANT: SCHNEIDER-MERGENER, JENS
TITLE OF INVENTION: METAL-BINDING CYSTEIN-FREE
TITLE OF INVENTION: THERAPEUTICAL PURPOSES,
TITLE OF INVENTION: METHODS FOR THEIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FRAGMENT TYPE:
ORIGINAL SOURCE:
                             TOPOLOGY: line MOLECULE TYPE: p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL:
ANTI-SENSE: I
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 7.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT:
                                                                             LENGTH: 17 amino acids
TYPE: amino acid
TYPE: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27.3%;
Local Similarity 37.5%;
es 6; Conservation
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6291639
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internal
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                                                                                                                                                                                                                     WBLT2010 (WO/US)
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SEQUENCE CHARACTERISTICS: LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
TYPOLOGY: linear
TOPOLOGY: linear
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                                                                                                                                                  RESULT 13
US-08-382-013A-9
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US-08-333-565-14
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                                                                                                                 Sequence 9, Application US/08382013A Patent No. 5631222
                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                    GENERAL INFORMATION:
APPLICANT: Shibat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: KORSME
                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/33,565
FILING DATE: 31-OCT-1994
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726A-000700
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Bcl-x/Bcl
TITLE OF INVENTION: REGULATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U
ZIP: 94301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08333565
Shibata, Kenji
Suzawa, Toshiyuki
Yamasaki, Motoo
Tanaka, Takeo
Tsukuda, Eiji
Yamada, Koji
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  : (415) --
(415) 326-2422
--- TD NO: 14:
                                                                                                                                                                                                                                                                                     Conservative
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/ENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
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                                                                                                                                                                                                                                                                                                  26.7%;
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                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                    Score 23; DB 1;
Pred. No. 2.5e+03;
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                                                                                                                                                                                                                                                                                                                  Length 17;
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NUMBER OF SEQUENCES: 7

CITLE OF INVENTION:

Endothelin-Antagonizing Peptide

Ohno, Tetsuj:

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RESULT 14
US-08-321-625-52
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: sinal-
TOPOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                                                                                     Sequence 52, Application US/08321625 Patent No. 5639860
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compactible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/382,013A
FILING DATE: 10-FEB-1995
ZIP: 22209
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: JP 155031/93
                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: ANTONELLI
                                                                                                                                                                                                                            APPLICANT:
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TELEPHONE: 212-819-8200
                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: JP 155;
FILING DATE: 25-UN-1993
RATTORNEY/AGENT INFORMATION:
NAME: Lippert, Nels T.
REGISTRATION NUMBER: 25888
                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                               APPLICANT:
                                        CITY: Arlington
STATE: VA
COUNTRY: USA
                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
STREET: 11
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                                                                                 CHLD.,

CHLD.,

CHLD.,

CHLD.,

TSUKUDA, Eiji

T: YOSHLDA, Mayumi

AT: BANDO, Chieko

ANT: YAMAGUCHI, Kazuo

ANT: YAMAGUCHI, Kazuo

ANT: MATSUDA, Yuzuru

ANTION: ENDOTHELIN-ANTAGONIZING PEPTIDE

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Y: U.S.A.
                                                                                      E: ANTONELLI, TERRY, STOUT & KRAUS
                                                                                                                                                                                                                                                      TANAKA, Takeo
MORISHITA, Yosh
MAKINO, Mika
CHIBA, Shigeru
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Floppy disk
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75.0%;
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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2.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Application US/08661479
Patent No. 5834209
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 75.0 Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                              FILING DATE: 11-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNWBER: US 08/333,565
FILING DATE: 31-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726A-000
TELEPHONE: (415) 326-2400
TELEPHONE: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 703-312-6666
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC compatible
COMPUTER: PM PC compatible
COMPUTER: PACENTIN Release #1.0, Version #1.25
SOFTWARE: Pacentin Release
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Bcl-x/Bcl
TITLE OF INVENTION: REGULATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 50
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-312-6600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: TERRY, David T.
REGISTRATION NUMBER: 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FRAGMENT TYPE: N-terminal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: both
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ENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
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Pred. No. 2.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 17;
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TYPE: amino acid
STRANDEDNESS: single
FOROLOGY: linear
MOLECULE TYPE: peptide
US-08-661-479-14

Query Match
Best Local Similarity 62.5%; Pred. No. 2.5e+03;
Matches 5; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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4 RSMGGKED 11
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Db
4 RQNGGWED 11
Search completed: June 13, 2006, 01:45:09

Job time: 49 secs
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Result
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                       seq length: 17 seq length: 17
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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                  US-10-289-454-376
US-10-346-190-157
US-10-428-254A-97
US-10-428-254A-92
US-10-289-456-157
US-10-328-2478A-10
US-10-622-124-100
US-10-622-087-70
US-11-037-396-62
US-11-037-396-62
US-09-988-493-251
US-09-988-493-251
US-10-481-261-225
US-10-834-397-225
US-10-834-397-254
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US-10-801-99-7

US-10-291-241-85

US-10-356-257-295

US-09-853-253-14

US-09-798-116-23

US-10-921-371-14

US-10-921-371-14

US-10-349-507-15

US-10-349-507-15

US-10-356-257-263

US-10-356-257-263

US-10-356-257-263

US-10-356-257-243

US-10-356-257-243
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85, Appli
295, Appli
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14, Appli
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15, Appl
163, Appl
263, App
9, Appli
92, Appl
157, Appl
10, Appl
10, Appl
70, Appl
62, Appl
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sequence sasts, A		Sequence 570, App		Sequence 284, App	Sequence 226, App	Sequence 224, App	Sequence 222, App	Sequence 57, Appl	Sequence 196, App	Sequence 2, Appli	Sequence 269, App	Sequence 102, App	Sequence 557, App	Sequence 2250, Ap	Sequence 52, Appl	Sequence 12, Appl	Sequence 2, Appli

ALIGNMENTS

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APPLICANT: Dade Behring Marburg GmbH
ITILE OF INVENTION: Carbohydrate deficient transferrin (CD
ITILE OF INVENTION: antibodies, their preparation and use
FILE REFERENCE: 2002/B001
CURRENT APPLICATION NUMBER: US/10/612,162
CURRENT FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 10230550.1
PRIOR FILING DATE: 2002-07-05
PRIOR FILING DATE: 2002-07-05
PRIOR FILING DATE: 2003-05-19
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO
LENGTH: 17
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US-10-801-990-7
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US-10-612-162-1
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US-10-612-162-1
                                                                                                                                                                                                                                                                                          Sequence 7, Application US/10801990 Publication No. US20050048574A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10612162 Publication No. US20040014145A1 GENERAL INFORMATION:
NUMBER OF SEQ ID NOS: 395
SOFTWARE: PatentIn version 3.2
SEQ ID NO 7
LENGTH: 17
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                                                                                FILE REFERENCE: SURR.121
CURRENT APPLICATION NUMBER: US/10/801,990
CURRENT FILING DATE: 2004-03-15
PRIOR APPLICATION NUMBER: US 60/455,037
PRIOR FILING DATE: 2003-03-14
                                                                                                                                                                                                   APPLICANT: Kantor, Aaron B.
APPLICANT: Schulman, Howard
APPLICANT: Becker, Christopher
TITLE OF INVENTION: BIOMARKERS FOR RHEUMATOID ARTHRITIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 17
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VVARSMGGKEDLIWELL 17
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APPLICANT: Robert Kendall Morrison
APPLICANT: Pia M. Challita-Eid
ITITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
ITITLE OF INVENTION: ENTITLED 161P2F10B USEFUL IN TREATMENT ANI
ITITLE OF INVENTION: CANCER
FILE REFERENCE: 51158-20062.20
CURRENT APPLICATION NUMBER: US/10/291,241
CURRENT APPLICATION NUMBER: US 10/005,480
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 0/062,109
PRIOR APPLICATION NUMBER: US 0/062,109
PRIOR APPLICATION NUMBER: US 60/282,739
PRIOR APPLICATION NUMBER: US 60/282,739
PRIOR APPLICATION NUMBER: US 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 103
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                  US-10-356-257-295
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; ORGANISM: Homo Sapiens
US-10-291-241-85
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                                     APPLICANT: LAZARUS, ROBERT A.
APPLICANT: MAUN, HENRY R.
APPLICANT: MAUN, HENRY R.
TITLE OF INVENTION: FVIIA Antagonists
FILE REFERENCE: P1950R1
CURRENT APPLICATION NUMBER: US/10/356,257
CURRENT FILING DATE: 2003-01-30
PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR FILING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 355
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                                                                                                                                                                                                                                  Sequence 295, Application US/10356257 Publication No. US20040087767A1 GENERAL INFORMATION:
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Matches
SEQ ID NO 295
LENGTH: 17
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APPLICANT: Agensys, Inc.
APPLICANT: Aya Jakobovits
APPLICANT: Arthur B. Raitano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity 50.0%;
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Local Similarity 100.0%; les 8; Conservative 0;
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Pred. No. 6.8e+02;
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CURRENT APPLICATION NUMBER: US/09/853,253
CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: 60/203,300
PRIOR FILING DATE: 2000-05-1
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 14
LENGTH: 17
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Best Local Similarity
Matches 3; Conserve
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US-09-853-253-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-853-253-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-798-116-23
                                     US-09-798-116-23
                                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: Verhagen, Anne Marie
APPLICATION INMERR: US/09/798,116
CURRENT FILING DATE: 2001-03-02
PRIOR FILING DATE: 2000-03-02
PRIOR FILING DATE: 2000-03-02
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Best Local Similarity
"hes 6; Conserv
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                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: JASPERS, STEPHEN
APPLICANT: SHEPPARD, PAUL
APPLICANT: DEISHER, THERESA
APPLICANT: BISHOP, PAUL
TITLE OF INVENTION: Z81g33-like Peptides
FILE REFERENCE: 00-30
                                              NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: M is methionine sulfoxide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial sequence FEATURE:
                                                                                                             ORGANISM: synthetic FEATURE:
                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 EDLIWE 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 QDILWE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 GGKEDLI-W 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09853253
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Score 28;
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Pred. No. 1.5e+03;
3; Mismatches (
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Pred. No. 1
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Length 17;
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Query Match

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US-10-921-371-14
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US-10-921-371-14
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                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/349,507
CURRENT FILING DATE: 2003-01-22
PRIOR APPLICATION NUMBER: 10/312,187
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: PCT/CA01/00913
PRIOR FILING DATE: 2001-06-20
PRIOR APPLICATION NUMBER: 60/213,174
PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/213,174
PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/254,932
PRIOR FILING DATE: 2000-12-13
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                                          SEQ ID NO 15
LENGTH: 17
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: JASPERS, STEPHEN
APPLICANT: SHEPPARD, PAUL
APPLICANT: DEISHER, THERESS
APPLICANT: BISHOP, PAUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15, Application US/10349507
Publication No. US20030199002A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 50.0 Matches 3; Conservative
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ORGANISM: Homo sapiens -10-349-507-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2004-08-19
PRIOR APPLICATION NUMBER: US/09/853,253
PRIOR FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: 60/203,300
PRIOR APPLICATION NUMBER: 60/203,300
PRIOR FILING DATE: 2000-05-11
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McCright, Brenton
APPLICANT: Lakowski , Bernard
TITLE OF INVENTION: CLK-2 NUCLEIC ACIDS, POLYPEPTIDES, AND USES THEREOF
FILE REFERENCE: 11202-006-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Hekimi , Siegfried
APPLICANT: Jiang , Ning
APPLICANT: Benard, Claire
APPLICANT: Kebir , Hania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: SHEPPARD, PAUL
APPLICANT: DEISHER, THERESA
APPLICANT: BISHOP, PAUL
TITLE OF INVENTION: ZSig33-like Peptides
FILE REFERENCE: 00-30
                                                                                                         NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn versi
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CURRENT FILING DATE: 2004-08-19
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TYPE: PRT
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50.0%;
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Pred. No. 1.5e+03;
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                           US-10-356-257-263
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US-10-267-565-15
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SEQ ID NO 15
LENGTH: 17
                                                                                                                                                Sequence 263, Application US/10356257
Publication No. US20040087767A1
GENERAL INFORMATION:
APPLICANT: LAZARUS, ROBERT A.
APPLICANT: MAUN, HENRY R.
TITLE OF INVENTION: EVIIA Antagonists
CURRENT APPLICATION NUMBER: US/10/356,257
CURRENT FILING DATE: 2003-01-30
PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR FILING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 355
SEQ ID NO 263
LENGTH: 17
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PRIOR APPLICATION NUMBER: 07/455,708
PRIOR FILING DATE: 1989-12-22
NUMBER OF SEQ ID NOS: 43
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CURRENT FILING DATE: 2003-01-23
PRIOR APPLICATION NUMBER: 09/401,839
PRIOR FILING DATE: 1999-09-22
PRIOR APPLICATION NUMBER: 08/459,151
PRIOR FILING DATE: 1995-06-02
PRIOR FILING DATE: 1995-06-02
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APPLICANT: Gubler, Ulrich
APPLICANT: Hulmes, Jeffery
APPLICANT: Podlaski, Frank
APPLICANT: Stern, Alvin
                                                                                                                                        FILE REFERENCE: P1950R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 08/205,011
PRIOR FILING DATE: 1994-03-02
PRIOR APPLICATION NUMBER: 07/857,023
PRIOR FILING DATE: 1992-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: PURIFICATION AND CHARACTERIZATION TITLE OF INVENTION: LYMPHOCYTE MATURATION FACTOR AND TITLE OF INVENTION: ANTIBODIES THERETO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1990-08-27
PRIOR APPLICATION NUMBER: 07/520,935
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5; Conserv
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Pred. No. 2.2e
0; Mismatches
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Pred. No. 2.2e+03;
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GENERAL INFORMATION:
APPLICANT: LAZARUS, ROBERT A.
APPLICANT: MAUN, HENRY R.
APPLICANT: MAUN, HENRY R.
ITITLE OF INVENTION: FVIIA Antagonists
FILE REFERENCE: P1950R1
CURRENT APPLICATION NUMBER: US/10/356,257
CURRENT FILING DATE: 2003-01-30
PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR FILING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 355
SEQ ID NO 243
LENGTH: 17
TYPE: PRT
US-10-356-257-243
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                                                                                                                                                                                                                                                                                                                                                     US-10-356-257-243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: peptide from nef pool 1 US-11-008-653-145
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                                                                                                                                                                                                                                                                                                                Sequence 243, Application US/10356257 Publication No. US20040087767A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local
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PRIOR FILING DATE: 2003-10-27
PRIOR APPLICATION NUMBER: GB 0322637.0
PRIOR FILING DATE: 2003-09-26
PRIOR APPLICATION NUMBER: GB 0322402.9
PRIOR FILING DATE: 2003-09-24
NUMBER OF SEQ ID NOS: 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/11/008,653
CURRENT FILING DATE: 2004-12-09
PRIOR APPLICATION NUMBER: PCT/GB2004/004038
PRIOR FILING DATE: 2004-09-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Schneider, Joerg
TITLE OF INVENTION: HIV Pharmaccines
FILE REFERENCE: 3742.1001-000
                   OTHER INFORMATION: sequence is synthesized
                                  ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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    Conservative

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Pred. No. 3.3e+03;
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Pred. No. 3.3e+03;
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                                                                                                                                                                  FILE REFERENCE: 1700.0360001

CURRENT APPLICATION NUMBER: US/10/289,454

CURRENT TELLING DATE: 2003-02-10

PRIOR APPLICATION NUMBER: US 60/396,636

PRIOR PILING DATE: 2002-07-19

PRIOR APPLICATION NUMBER: PCT/1802/00166

PRIOR FILING DATE: 2002-01-21

PRIOR APPLICATION NUMBER: US 10/050,902

PRIOR APPLICATION NUMBER: US 60/331,045

PRIOR FILING DATE: 2001-11-07

PRIOR FILING DATE: 2001-11-07
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; OTHER INFORMATION: CepsilonH3 foreward US-10-289-454-376
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SEQ ID NO 4
LENGTH: 17
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                                                                                                              SOFTWARE: PatentIn version 3.2 SEQ ID NO 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 376, Application US/10289454 Publication No. US20030157479A1
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CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: JP 2001-131688
PRIOR FILING DATE: 2001-04-27
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TITLE OF INVENTION: ADENOVIRUS VECTOR
FILE REFERENCE: 081356/0163
                                                                                                                                                    NUMBER OF SEQ ID NOS: 386
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jennings, Gary APPLICANT: Sonderegger, I
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APPLICANT: HAYAKAWA, TAKAO
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bachmann, Martin
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                                  TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 4e+03;
2; Mismatches
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APPLICANT: Maurer, Patrick
APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
CURERI FILIR CONTONIONER: Wolfgang A.
FILE REFERENCE: 1700.0290003
CURRENT FILING DATE: 2003-01-17
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
SOFTWARE: Patrentin version 3.1
SEQ ID NO 157
LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
COTURE INFORMATION: Oliconological de crimos
Search completed: June 13, 2006, 02:00:46 Job time : 67 secs
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is derived by analysis of the total score distribution.
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US-11-259-402-83
US-11-539-402-83
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US-11-508-063A-18
US-11-508-063A-18
US-10-508-063A-18
US-10-508-1300-62
US-10-981-300-67
US-11-134-445-68
US-11-122-986-435
US-11-122-986-435
US-11-123-610-172
US-10-984-473-26
US-10-984-473-26
US-10-984-473-27
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ALIGNMENTS

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APPLICANT: Devaux, Brigitte
APPLICANT: Devaux, Brigitte
APPLICANT: Devaux, Brigitte
APPLICANT: Eaton, Dan L.
APPLICANT: Hass, Philip E.
APPLICANT: Judice, J. Kevin
APPLICANT: Kirchhofer, Daniel
APPLICANT: Suggett, Shelley
TITLE OF INVENTION: Human Anti-Factor IX/IXa Antibodies
FILE REFERENCE: P1661R2
CURRENT APPLICATION NUMBER: US/10/614,959
CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/383,667
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/383,667
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/383,667
PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1999-08-26
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/098,233
PRIOR APPL
                                                                                                                                                                                                                                                  Sequence 21, Application US/11251734 Publication No. US20060100134A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 3; Conserv
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                                                                                         APPLICANT: AVENTIS PHARMACEUTICALS INC
APPLICANT: GUO, Yong
APPLICANT: MORSE, Clarence C
APPLICANT: GUO, Yong
APPLICANT: MORSE, CLARENCE C
APPLICANT: YAO, Zhengbin
TITLE OF INVENTION: MEMBRANE PENETRATING PEPTIDES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION
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Pred. No. 1.2e+03;
2; Mismatches 0;
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; FEATURE:
; OTHER INFORMATION: Synthetic peptide
US-11-251-734-21
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; ORGANISM: human
US-10-539-402-81
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CURRENT FILING DATE: 2005-06-17
PRIOR APPLICATION NUMBER: US 60/435,893
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: EP 03000615
PRIOR APPLICATION NUMBER: EP 03000615
PRIOR FILING DATE: 2003-01-15
NUMBER OF SEQ ID NOS: 108
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-539-402-81
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CURRENT FILING DATE: 2005-10-17
PRIOR APPLICATION NUMBER: US 09/933,780
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR FILING DATE: 2000-08-25
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                      Sequence 5, Application US/11039005 Publication No. US20060115856A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 81, Application US/10539402 Publication No. US20060115477A1 GENERAL INFORMATION:
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SEQ ID NO 21
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Best Local Similarity
APPLICANT: Goldknopf, Ira L.
APPLICANT: Sheta, Essam A.
APPLICANT: Sheta, Essam A.
APPLICANT: Appel, Stanley H.
APPLICANT: Simpson, Ericka P.
APPLICANT: Simpson, Ericka P.
APPLICANT: Yen, Albert A.
TITLE OF INVENTION: AN 2'-5'-OLIGOADENYLATE SYNTHETASE LIKE PROTEIN AS A BIOMARKER
TITLE OF INVENTION: FOR NEURODEGENERATIVE DISEASE
FILE REFERENCE: PROT-P0013US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Xerion Pharmaceuticals AG
APPLICANT: Tufts University
TITLE OF INVENTION: Neuropilin-1 Inhibitor
FILE REFERENCE: XE12EPC
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Pred. No. 1.2e+03;
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APPLICATION INTERPOLATION SPECIFICITY EXCHANGERS THAT REDIRECT TITLE OF INVENTION: SPECIFICITY EXCHANGERS THAT REDIRECT TITLE OF INVENTION: ANTIBODIES TO A PATHOGEN
FILE REFERENCE: TRIPER: 7AUC4CP1C
CURRENT APPLICATION NUMBER: US/11/121,282
CURRENT FILING DATE: 2005-05-03
PRIOR APPLICATION NUMBER: 10/372,735
PRIOR FILING DATE: 2003-02-21
PRIOR FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: 10/234,579
PRIOR APPLICATION NUMBER: 10/234,579
PRIOR APPLICATION NUMBER: 09/839,666
PRIOR FILING DATE: 2001-04-19
PRIOR APPLICATION NUMBER: 09/839,666
PRIOR APPLICATION NUMBER: 09/839,666
PRIOR APPLICATION NUMBER: 09/832,106
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Best Local Similarity
Matches 3; Conservi
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Best Local Similarity 80.0
                                                                                                                                                                                                                                                           Sequence 52, Application US/11121282
Publication No. US20060094862A1
GENERAL INFORMATION:
APPLICANT: Matti Sallberg
APPLICANT: Jan-Ingmar Flock
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 83
LENGTH: 17
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APPLICANT: Xerion Pharmaceut
APPLICANT: Tufts University
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 5
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Publication No. US20060115477A1
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CURRENT APPLICATION NUMBER: US/10/539,402
CURRENT FILING DATE: 2005-06-17
PRIOR APPLICATION NUMBER: US 60/435,893
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: EP 03000615
PRIOR FILING DATE: 2003-01-15
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CURRENT FILING DATE: 2005-01-19
PRIOR APPLICATION NUMBER: 60/632,218
PRIOR FILING DATE: 2004-12-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Neuropilin-1 Inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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100.0%; Pred. No. 3.7e+03;
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Pred. No.
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. 1.7e+03;
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TITLE OF INVENTION: DITECTION OF ANALYTES
FILE REFERENCE: 07257/042001
CURRENT APPLICATION UNMBER: US/11/334,622
CURRENT FILING DATE: 2006-01-17
PRIOR APPLICATION NUMBER: US/9/554,000
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 1997-03-14
NUMBER: 05/818,252
PRIOR FILING DATE: 1997-03-14
NUMBER: 0F SEQ ID NO5: 56
SOFTWARE: FASHSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 17
Sequence 18. Application US/10508063A
Publication No. US20060088832A1
GENERAL INFORMATION:
APPLICANT: Deutsches Krebsforschungszentrum
APPLICANT: ANGEL, Peter
APPLICANT: HRSTENBERGER, Gerhard
APPLICANT: RICHTENBACH, Ute
APPLICANT: RICHTER, Harmut
APPLICANT: HESS, Jochen
APPLICANT: HESS, Jochen
TITILE OF INVENTION: TAP-70, A NOVEL MARKER FOR
FILLE REFERENCE: 31304-701.831
CURRENT APPLICATION NUMBER: US/10/508,063A
CURRENT FILING DATE: 2004-09-16
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US-11-334-622-12
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                                                                                                                                                                                                                                                                                              US-10-508-063A-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Gallus gallus US-11-334-622-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 19.8%;
Best Local Similarity 66.7%;
Matches 2; Conservative
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Best Local Similarity 50.0%;
Matches 3; Conservative
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PRIOR FILING DATE: 2000-09-19
PRIOR APPLICATION NUMBER: 09/246,258
PRIOR FILING DATE: 1999-02-08
NUMBER OF SEQ ID NOS: 199
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APPLICANT: Miyawaki,
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                    A NOVEL MARKER FOR
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Pred. No. 3.7e+03;
1; Mismatches 0
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Pred. No. 3.7e+03;
                                                                      EPITHELIAL TUMORS
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APPLICANT: GIORGIO SENALDI
APPLICANT: GADI GAZIT-BORNSTEIN
TITLE OF INVENTION: ANTI-PROBERDIN ANTIBODIES, AN
TITLE OF INVENTION: FOR MAKING AND USING THE SAM
FILE REFERENCE: ABGX-005
CURRENT APPLICATION UNMBER: US/10/981,300
CURRENT FILING DATE: 2004-11-03
NUMBER OF SEQ ID NOS: 71
SOPTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 63
LENGTH: 17
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US-10-981-300-62
J Sequence 62, Application US/10981300
Publication No. US20060093599A1
Publication No. US20060093599A1
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; TYPE: PRT
; ORGANISM: Artificial
; FEATURE;
; FEATURE;
; OTHER INFORMATION: Synthetic peptide
US-10-508-063A-18
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Best Local Similarity
Thes 3; Conserve
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US-10-981-300-63
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                                                                                                US-10-981-300-63
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APPLICANT: GADI GAZIT-BORNSTEIN
TITLE OF INVENTION: ANTI-PROBERDIN ANTIBODIES, AND METHODS
TITLE OF INVENTION: FOR MAKING AND USING THE SAME
FILE REFERENCE: ABGX-005
CURRENT APPLICATION NUMBER: US/10/981,300
CURRENT FILING DATE: 2004-11-03
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 62
LENGTH: 17
                                                                                                                                                                                                                                                                                                                      Sequence 63, Application US/10981300 Publication No. US20060093599A1 GENERAL INFORMATION:
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                              Best Local Similarity Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                              Query Match
                                                                                                          LENGTH: 17
TYPE: PRT
ORGANISM: Homo sapiens
12 LIW 14
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ilarity 37.5%;
Conservative
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                                               18.6%;
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                              Score 16; DB 6;
Pred. No. 5.3e+03;
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Pred. No. 5.3e+03
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Pred. No. 5.3e+03;
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; OTHER INFORMATION: substituted CDR
US-10-544-050-67
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US-10-544-050-67
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CURRENT FILING DATE: 2005-05-23
PRIOR APPLICATION NUMBER: 10/792,893
PRIOR FILING DATE: 2004-03-05
PRIOR PELICATION NUMBER: 10/383,614
PRIOR PELICATION NUMBER: 10/277,969
PRIOR APPLICATION NUMBER: 10/277,969
PRIOR APPLICATION NUMBER: 10/277,969
PRIOR APPLICATION NUMBER: 10/270,148
PRIOR APPLICATION NUMBER: 10/200,148
PRIOR APPLICATION NUMBER: 10/200,148
PRIOR PILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: 10/141,647
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US-11-134-445-68
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SEQ ID NO 67
LENGTH: 17
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Publication No. US20060094677A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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Publication No.
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Best Local
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CURRENT APPLICATION NUMBER: US/10/544,050
CURRENT FILING DATE: 2005-07-29
PRIOR APPLICATION NUMBER: 60/446380
PRIOR FILING DATE: 2003-02-10
NUMBER OF SEQ ID NOS: 76
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APPLICANT: HEIKKILA, ELIZABETH MARGARET
APPLICANT: SENCHYNA, DIANE MICHELLE
APPLICANT: FLANAGAN, JOHN GERARD
APPLICANT: THOMBSON, JOHN E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Davies Julian
TITLE OF INVENTION: Abeta Binding Molecules
                                            PRIOR FILING DATE: 2004-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: INHIBITION OF APOPTOSIS-SPECIFIC eIF-5A ("eIF-5A1") WITH TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES AND SIRNAS AS ANTI-INFLAMMATCHIE OF INVENTION: THERAPEUTICS FILE REFERENCE: 10799/125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: TAYLOR,
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ORGANISM: Artificial sequence
                                                                                                                               OR FILING DATE: 2002-05-07
OR APPLICATION NUMBER: 09/909,796
OR FILING DATE: 2001-07-23
OR APPLICATION NUMBER: 60/476,194
OR FILING DATE: 2003-06-06
OR APPLICATION NUMBER: 60/504,731
OR FILING DATE: 2003-09-22
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les 3; Conserv
                                                               APPLICATION NUMBER: 60/528,249 FILING DATE: 2003-12-10 APPLICATION NUMBER: 60/557,671
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Prior Application data removed SEQ ID NOS: 95
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100.0%; Pred. No. 5.3e+03
cive 0; Mismatches 0
                     See File Wrapper or
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SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 435
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SEQ ID NO 68
LENGTH: 17
TYPE: PRT
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                                                                    Matches
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CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
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                                                                                                                                                                        LENGTH: 17
TYPE: PRT
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PRIOR FILING DATE: 2002-11-06
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PRIOR APPLICATION NUMBER: 60/424,370
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PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,376
PRIOR FILING DATE: 2002-11-06
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PRIOR FILING DATE: 2002-11-05
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APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
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                                                                                                                                                                                                                                                                                                                FILING DATE:
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Pred. No. 5.3e+03;
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5.3e+03;
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WESULT 14 US-11-23-610-172 ; Sequence 172, Application US/11223610 ; Publication No. US20060111289A1

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Sequence 39 Application US/11334622
Publication No. US20060112440A1
GENERAL INFORMATION:
APPLICANT: Tsien, Roger Y.
APPLICANT: Tsien, Roger Y.
TITLE OF INVENTION: FLUORESCENT PROTEIN SENSORS FOR TITLE OF INVENTION: DETECTION OF ANALYTES
FILE REFERENCE: 07257/042001
CURRENT APPLICATION NUMBER: US/11/334,622
CURRENT APPLICATION NUMBER: US/09/554,000
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: 08/818,252
PRIOR FILING DATE: 1997-03-14
NUMBER OF SEQ ID NOS: 56
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 39
LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
COTHER INFORMATION: Calmodulin binding peptide-2
US-11-334-622-39
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US-11-334-622-39
Search completed: June 13, 2006, 02:00:58 Job time : 7 secs
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FILE REFERENCE: HOLLAND-09516
CURRENT APPLICATION NUMBER: US/11/223,610
CURRENT FILING DATE: 2005-09-09
NUMBER OF SEQ ID NOS: 181
SOFTWARE: Patentin version 3.3
SEQ ID NO 172
LENGTH: 17
TYPE: PRT
                                                                                                                                                               Query Match 18.6%; Score 16; DB 7; Length 17; Best Local Similarity 40.0%; Pred. No. 5.3e+03; Matches 2; Conservative 2; Mismatches 1; Indels
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APPLICANT: Mizejewski, Gerald J.
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Local Similarity 60.0%;
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ABM83783
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ADL24413
ADU17885
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110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	110	
95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	95.7	
698	698	869	869	698	869	869	698	869	869	869	679	679	679	679	679	679	679	679	679	679	
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ADK15868	ADH89359	ADH34559	ADF74800	ADF74796	ADD45282	ABP72819	AAY50717	AAW54354	AAR66492	AAR12499	AEE47501	ADY57103	ADY57102	ADY54149	ADT55066	ADU00630	ADU17468	ADU17611	ADT98423	ADT36538	
Adk15868	Adh89359	Adh34559	Adf74800	Adf74796	Add45282	Abp72819	Aay50717	Aaw54354	Aar66492	Aar12499	Aee47501	Ady57103	Ady57102	Ady54149	Adt55066	Adu00630	Adu17468	Adu17611	Adt 98423	Adt36538	
Human t	Human t	Human t	Human N	Human N	Human P	Human t	Human ser	Serotrans	Human tra	Human t	Aee47501 Human tra	Human s	Human s	Amino a		Human t	Human t	Human t	Human t	Partial	
tra	tra	tra	NOV	VOV	Pro	tra	er	91	ra	tra	tra	ser	ser	acı	par	tra	tra	tra	tra	ב	,

ALIGNMENTS

RESULT 1 ADG446076 ID ADG44 XX ADG4 AX ADG4 XX ADG4 XX ADG4 XX Anti XX Anti XX Anti XX NON. XX HOMC XX ALTI XX ALTI XX HOMC XX HOMC XX ALTI XX HOMC XX antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. Human CDT peptide fragment ADG46076; 05-JUL-2002; 2002DE-01030550. 19-MAY-2003; 2003EP-00011334. 07-JAN-2004. EP1378521-A1. Homo sapiens 25-MAR-2004 ADG46076 standard; peptide; (DADE-) DADE BEHRING MARBURG GMBH (first entry) 23 B

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSMCGKEDLIWELL, TTEDSIAKIMNGRADAMSLDGGF, SKLSMGSGLNISSEN and YEKYLGEEYVAAV (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization. Claim 4; SEQ ID NO 2; 21pp; German.

WPI; 2004-073743/08.

Althaus H;

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RESULT 2
AAB84371
ID
AAB84371

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Best Local S
Matches 23
                                             The present sequence represents human transferrin polypeptide. Transferrin polypeptides and polynucleotides are used to prevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. erebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sumburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          production of antibodies that specifically bind CDT from the aqueous phase. The antibodies are useful for diagnosis of alcoholism, CDT, whice lacks glycosylation at Asn413 and/or Asn 611, is present only in alcoholics. The antibodies allow direct detection of CDT in solution, eliminating the need for immobilising it on a solid phase (as required when using known antibodies). ADG46075-ADG46094 represent CDT peptide fragments used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; transferrin; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; disorder lischemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; epithelial cell proliferation; aging; organ transplant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; Page 286-287; 290pp; English.
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                        preservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecule encoding a human tranferrin protein is used in preventing, treating or ameliorating a medical condition.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001-381910/40
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                        polypeptides can also
increase or decrease
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요
                    storage
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Matches 22
                                              The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants granulocyte colony stimulating factor receptor, glucagon, interleukin 6 platelet-derived endothelial cell growth factor, cyclin-dependent kinas inhibitor 1C, cellular tumour antigen P53, and vasoactive intestinal polypeptide receptor 2. The polypeptides and their associated mucleic acids are useful for identification of variant sequences and detection candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various disorders including cardiovascular diseases such as arteriosclerosis, myocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as hypertrophy, immune disorders such as immune complex nephritis, multiple sclerosis such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Angiotensin converting enzyme splice variant; ACEV; interleukin 6; granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P5; cyclin-dependent kinase inhibitor 10; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; myocardial infarction; coronary arterial thrombosis; renal disease; diabetic nephropathy; muscular disease; immune disorder; sarcoidosis; multiple sclerosis; immune complex nephritis; deep vein thrombosis; nonarcoidocic pulmonary granulomatous disease; endothelial abnormality;
                                                                                                                                                                                                                                                                                                                                  Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibo
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                cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases as asbestosis and vascular pathologies involving an endothelial
                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; AAS06080
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10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vascular disorder; asbestosis.
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                                                                                                                                                                                                                                                                                                   4; Fig 80; 519pp; English.
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22; Conser
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99IL-00133455.
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95.7%;
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Pred. No. 6.8e-09;
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polynucleotides. The human MEPR polypeptides, polynucleotides and modulators are useful for diagnosing, treating or preventing disorders associated with aberrant expression of MEPR, particularly cell proliferative disorders (e.g. arteriosclerosis, atherosclerosis, properties, particularly cell proliferative disorders (e.g. arteriosclerosis, atherosclerosis, polycythemia cirrhosis, hepatitis, paroxysmal nocturnal hemoglobinuria, polycythemia vera, psoriasis, primary thromobocytopenia or cancer), developmental disorders (e.g. renal tubular acidosis, anemia or mental retardation), neurological disorders (e.g. Alzheimer's disease, Parkinson's disease or epilepsy), autoimmune/inflammatory disorders (e.g. AIDS, allergies, asthma, autoimmune thyroiditis, contact dermatitis, Crohn's disease, diabetes mellitus, glomerulonephritis, Goodpasture's syndrome, gout, Graves' disease, Hashimoto's thyroiditis, irritable bowel syndrome,
                                                                                                                                                                                                                                                              allergy or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-JAN-2002; 2002US-0348769P.
18-JAN-2002; 2002US-035070H.
19-MAR-2002; 2002US-0366059P.
10-MAY-2002; 2002US-0379907P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antianemic; antiparkinsonian; nootropic; anticonvulsant; cytostatic; antiatreriosclerotic; antiasthmatic; immunosuppressive; antithyroid; hepatotropic; dermatological; antidiabetic; nephrotropic; antigout; thyromimetic; neuroprotective; osteopathic; antiarthritic; uropathic; antiparasitic; antihelmintic; antipsoriatic; ophthalmological; virucide; antiphematic; haemostatic; antibacterial; protozoacide; fungicide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JAN-2003; 2003WO-US001485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JUL-2003
                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                           New human metalloproteins and polynucleotides, useful for diagnosing,
                                                                                                                                                                                                                                                                                                                                                                                                           Emerling
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEPR; metalloprotein; anti-HIV; antiallergic; antiinflammatory; human;
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                                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy; transgenic.
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DB; ACF35812.
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                                                                                                                                                                                                                              1; Page 146-148; 153pp; English
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                                                                                                                                                                                                                                                                            or anemia), multiple sclerosis,
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M, Chien D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                              relates to novel human metalloproteins (MEPR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MEPR)
                                                                                                                                                                                                                                                                                                                                                                                                         Gorvad
Jin P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptide (Id-7509328CD1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 110; DB 4;
Pred. No. 1.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                         AE, Becha SD, I
Chawla NK, Yue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                              osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
                                                                                                                                                                                                                                                                                                                                                                                                       Richardson TW;
e H, Khare R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                              cancer
                                                                                                                                                                                                and encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                         Marquis JP;
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The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders, endocrine autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide

Claim 27;

Page; 190pp; English.

molecules, e.g.

w diagnostic and therapeutic polynucleotides and polypeptides, use diagnosing a condition, disease or disorder associated with human lecules, e.g. autoimmune or inflammatory disorders, in gene therap

therapy or

in gene mapping.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
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Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy T. Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gersi Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve J Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kil Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-NOV-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM83782;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2003; 2003WO-US028227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004023973-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2004-329368/30.
DB; ACN42434.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 575 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTEDCIAKIMNGEADAMSLDGGF 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and therapeutic pprotein SEQ ID NO:4031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and therapeutic polynucleotide; dithp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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.3e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Au AP, Gerstin EH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SC, Reddy TP;
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Best Local S
Matches 22
The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders, autoimmune/inflammatory disorder, developmental disorders, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline
                                                                                                                                                                                                                                                                                                                                                                                                                                           Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES,
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietz,
                                                                                                                                                                                                                                                                                              New diagnostic and therapeutic polynucleotides and polypeptides, use in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                         Claim 27; Page; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; ACN42433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human diagnostic and therapeutic pprotein SEQ ID NO:4030.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM83781;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM83781 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 609
                                                                                                                                                                                                                                                                             in gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INCY-) INCYTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                                                                       2004-329368/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        321 TTEDCIAKIMNGEADAMSLDGGF 343
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                                                                                                                                                                                                                                                                                                                                                                                                                           Kwong M, I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                             Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95.7%;
95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
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Pred. No. 1.4e-08;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polynucleotide; dithp
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Matches 22
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Peralta CH, Anderso
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Peralta CH, Spiro P
Peralta CH, North
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Xu Y, Kwong M, Po
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with human molecules, e.g. cell proliferative disorders, autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying indiverse from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germ gene therapy. The present sequence represents a dithp protein of the
                                                                                                                                                                                                                                      The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy. The present sequence represents a dithp protein of invention. Note: The sequence data for this patent is not represe the printed specification, but was obtained in electronic format from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New diagnostic and therapeutic polynucleotides and polypeptides, use in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 27;
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12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in gene mapping.
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DB; ACN42431.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Page; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anderson SB, Rio
Spiro PA, Stewart
M, Policky JL, I
hi X, Suarez CJ;
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Suchorolski MT, Altus CM, Pitts
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                     The invention relates to novel human metal binding proteins (MBP) and CC encoding polynucleotides. The MBPs can be expressed by standard CC recombinant methodology. The polypeptides and polynucleotides are useful CC in diagnosing, treating and preventing diseases or conditions associated CC with the decreased expression or overexpression of MBP, such as cell CC proliferative (e.g. cancer, atherosclerosis), neurological (e.g. AIDS, CC epilepsy, Huntington's disease, stroke), immune/inflammatory (e.g. AIDS, CC allergies) and developmental (e.g. Hypothyroidism, Cushing's syndrome) CC disorders, or infections. These are also useful in assessing the effects of exogenous compounds on the expression of nucleic acid and amino acid CC sequences of MBP. The MBP or its fragments are useful in screening CC compounds for effectiveness as agonist or antagonist of the polypeptides, CC or in altering the expression of the target polynucleotide and compounds that specifically bind to or modulate the activity of the polypeptide. An CC microarray comprising different nucleotide molecules affixed at distinct CC physical locations on a solid substrate, where at least one mucleotide comprises a first oligonucleotide of polynucleotide sequence compounds to the present that a least 30 contiguous nucleotides of the present segments are presents a human MBP and expression for profellar The arresent segments are presented as human MBP and expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human metal binding proteins (MBP), useful for diagnosing, treating and preventing diseases or conditions associated with the aberrant MBP expression e.g. cancer, AIDS, epilepsy, or infections.
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antiallergic; antiinflammatory; thyromimetic; transgenic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 628 AA;
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   present sequence represents a human MBP
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Pred. No. 1.5e-08;
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          The invention relates to novel diagnostic and therapeutic polynucleotides considered from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be CC used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine CC autoimmune/inflammatory disorder, gastrointestinal disorders, or CC disorder, neurological disorders, gastrointestinal disorders, or CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide CC polymorphisms, as molecular weight markers, and for somatic or germline CC invention. Note: The sequence represents a dithp protein of the CC invention. Note: The sequence data for this patent is not represented in CC the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
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Peralta CH,
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                                                                                                                                                                                                                                                                                        Claim 27; Page; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; ACN42430.
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e TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV
Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin,
Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton GM, Policky JL, Hurwitz BL, Ma Y, Jackson JL, G:
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Pred. No. 1.
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                                                              The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polyapptides may be used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine autoimmune/inflammatory disorder, developmental disorder, endocrine CC disorder, neurological disorders, gastrointestinal disorders, or CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide CC polymorphisms, as molecular weight maxiers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the cc invention. Note: The sequence data for this patent is not represented in content of the printed specification, but was obtained in electronic format directly content in the protein of the content of the printed specification, but was obtained in electronic format directly content in the protein of the content of the content of the printed specification, but was obtained in electronic format directly content in the content of th
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N-PSDB; ACN42435.
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                                                                                     The sequence represents an angiotensin converting enzyme splice variant CC (ACEV) polypeptide. The polypeptides of the invention include variants of CC granulocyte colony stimulating factor receptor, glucagon, interleukin 6, CC platelet-derived endothelial cell growth factor, cyclin-dependent kinase cinhibitor 1C, cellular tumour antigen P53, and vasoactive intestinal CC polypeptide receptor 2. The polypeptides and their associated nucleic concide are useful for identification of variant sequences and detection of candidate compounds capable of binding the molecules. The sequences of CC the invention can be used in the treatment and diagnosis of various cisorders including cardiovascular diseases such as arteriosclerosis, CC myocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as hypertrophy, concern, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases such as asbestosis and vascular pathologies involving an endothelial
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Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Angiotensin converting enzyme splice variant; ACEV; interleukin 6; granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P53; cyclin-dependent kinase inhibitor 1C; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; myocardial infarction; coronary arterial thrombosis; renal disease; diabetto nephropathy; muscular disease; immune disorder; sarcoidosis; multiple sclerosis; immune complex nephritis; deep vein thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Levine Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-NOV-2000; 2000WO-IL000766.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Fig 38; 519pp;
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Pred. No. 1.5e-08;
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                           The invention relates to novel diagnostic and therapeutic polynucleotides content from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorder, endocrine disorder, neurological disorder, developmental disorder, endocrine disorder, neurological disorder, gastrointestinal disorder, endocrine disorders, or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                 New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV; Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH; Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES; Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietzer Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                     Claim
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12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human diagnostic and therapeutic pprotein SEQ ID NO:4029.
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DB; ACN42432.
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                                                                                                                                                                                                                                                                                                                                 27; Page; 190pp; English
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                                                                                                                                                                                                                                                                                                                                                                    mapping
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95.7%;
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Pred. No. 1.
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.5e-08;
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Sequence 646

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RESULT 13
AAU02937
ID AAU02937
XX AAU02937
AC AAU02
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                                                                                        The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants of granulocyte colony stimulating factor receptor, glucagon, interleukin 6, CC platelet-derived endothelial cell growth factor, cyclin-dependent kinase inhibitor 1C, cellular tumour antigen P3, and vasoactive intestinal colonypeptide receptor 2. The polypeptides and their associated nucleic colds are useful for identification of variant sequences and detection of candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various clisorders including cardiovascular diseases such as arteriosclerosis, compocardial infarction and coronary arterial thrombosis, renal diseases such as diabetic nephropathy, muscular diseases such as hypertrophy, cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases such as asbettosis and vascular pathologies involving an endothelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibodies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Fig 37; 519pp; English.
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10-DEC-1999;
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99IL-00133455
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                                                                       as deep vein
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Pred. No. 1.5e-08;
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DB 4; Length 665;

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RESULT 14
ABP72820
ID PP72
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Best Local Similarity
Matches 22; Conserv
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                      Binding-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiparkinsonian; nootropic; anti-HIV; antiasthmatic; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; transferrin; neuroprotective; cerebroprotective; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human transferrin (mature polypeptide).
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/note= "Iron binding site"
517
/note= "Iron binding site"
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                                                                                                                                                                                                                                                                                                                                                                     /note= "N-glycosylated"
425. .427
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                                                                                                                                                                                                                                                                                             /note= "Iron binding site"
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Pred. No. 1.6e-08;
0; Mismatches 1
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RESULT 15
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ID ADH89
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AC ADH89
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ADH89360 standard; protein; 679

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15-APR-2004 ADH89360;

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373 TTEDCIAKIMNGEADAMSLDGGF 395

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                                                                                          Matches
                                                                                                                        Query Match
Best Local (
                                                                                                                                                                                                                                                                                                        treating a disease or disease symptom, or for delivering a therapeutic agent complexed to the ferric iron of transferrin to the inside of a cell or across the blood-brain barrier. The modified fusion protein, or a nucleic acid encoding it, can be used in the diagnosis, prognosis, prevention and/or treatment of diseases and/or disorders of the endocrine, nervous, immune, respiratory, cardiovascular, reproductive and digestive systems, diseases and/or disorders relating to the blood or to cell proliferation, inflammatory conditions, and to treat viral, fungal, the province of the conditions of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New fusion protein, useful in the diagnosis and treatment of diseases disorders relating to the respiratory, cardiovascular and digestive systems, comprises a transferrin protein fused to a therapeutic protein
                                                                                                                                                                                                                     Sequence 679 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 280-281; 298pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-332916/31.
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30-NOV-2001; 2001US-0334059P
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                                                                                          22;
                                                                                                                           Similarity
TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                                                                                                                                                                                                 or parasitic
                                                                                             Conservative
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/note= "Hinge region"
585
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                                                                                                                                                                                                                                                                                    inflammatory conditions, tic infection
                                                                                                                           95.7%;
95.7%;
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                                                                                                                           Score 110; DB 6;
Pred. No. 1.6e-08;
                                                                                                Mismatches
                                                                                                                                                      Length 679
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protein

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Best Local S
Matches 22
                                                                                                                                                                                 This invention relates to a novel fusion protein which comprises a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region. The invention may be useful for the development of compounds with cytostatic, antibacterial, virucide, antiparasitic, immunosuppressive or antiarthritic activity. In addition, the sequences disclosed may be useful for gene therapy. The fusion protein is useful for preparing a composition for treating a disease or disease symptom in a patient for example septic shock, endotoxic shock, cachexia syndromes associated with bacterial, viral or parasitic infections, neoplasm, autoimmune disease, arthritis or adverse effects associated with treatment for preventing graft rejection. The present sequence is that of the mature human transferrin protein which is related
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New fusion protein comprising a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region, useful for preparing a composition for treating e.g., septic shock, neoplasm or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-AUG-2001; 2001US-0315745P.
30-NOV-2001; 2001US-03340599.
30-AUG-2002; 2002US-00231494.
30-AUG-2002; 2002US-0406977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fusion protein; transferrin protein; glycosylation; antibody variable region; cytostatic; antibacterial; virucide; antibarastit; immunosuppressive; antiarthritic; gene therapy; septic shock; endotoxic shock; cachexia syndrome; bacterial in
                                                                                                                                                                      ç
                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 3; 82pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             viral infection; parasitic infection; neoplasm; autoimmune disease; arthritis; graft rejection; human.
                                                                                                                                     Sequence 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human transferrin protein mature amino acid sequence
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                                                                                                                                                                      the invention.
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 373
                                                                                   Similarity
TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prior CP,
                                                                   Conservative
                                                                                                                                     ΑĄ,
                                                                                 95.7%;
95.7%;
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                                                                                   Score 110; DB 8; Length 679; Pred. No. 1.6e-08;
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Search completed: June 13, 2006, 03:58:17 Job time : 149.824 secs

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Result
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Maximum DB seq
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 Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being p
                                                                                                                                                                                                                                                                                                                       Score
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Maximum Match 100%
Listing first 45 summaries
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                                       AE0819
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lactotransferrin p
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                                                                                                                                                                                                                                                     transferrin - mous
transferrin - pig
                                                                                                                                                                                                                                                                                                                      Description
                             conserved hypothet
                                                                         virulence
                                                                               virulence protein
                                                                                             probable membrane
                                                                                                       transferrin - bovi
                                                                                                                 probable membrane-
                                                                                                                                     phosphate ABC
                                                                                                                                                transferrin
                                                                                                                                                         transferrin precur
                                                                                                                                                                                                                                 lactoferrin precur
transferrin precur
outer capsid prote
                                          hypothetical prote
                                                                                                                           hypothetical
                                                                                                                                                                                          lactoferrin
                                                                                                                                                                                                                                                                            transferrin
                                                                                                                                                                                                                                                                                                 transferrin precur
                                                                                                                                                                                                                        .actotransferrin p
                                                                                                                                                                                                                                                                                                                                                                                   printed,
                                                            sensor p
                                                                                                                                                                                                                                                                           precur
                                                                                                                           prote
                                                                                                                                      tran
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ALIGNMENTS

A;Molecule type: DNA
A;Residues: 1-72;291-300 <ADR>
A;Residues: 1-72;291-300 <ADR>
A;Residues: 1-72;291-300 <ADR>
A;Cross-references: UNIPARC:UPI000016B0AC; UNIPARC:UPI000016B0AD; EMBL:M15673
A;Cross-references: UNIPARC:UPI000016B0AC; UNIPARC:UPI000016B0AD; EMBL:M15673
A;Cross-references: UNIPARC:UPI000016B0AD; EMBL:M15673
Biochem. Biophys. Res. Commun. 119, 273-281, 1984
A;Title: Molecular cloning and sequence analysis of cDNA for human transferrin.
A;Reference number: A32739; MUID:84153910; PMID:6322780
A;Accession: A32739 A;Molecule type: DNA
A;Residues: 73-263,'E',265-328,'N',330-562 <PAR>
A;Residues: 73-263,'E',265-328,'N',330-562 <PAR>
A;Residues: 73-263,'E',265-328,'N',330-562 <PAR>
A;Cross-references: UNIPARC:UPI000174484; EMBL.M1361
A;Adrian, G.S.; Korinek, B.W.; Bowman, B.H.; Yang, F.
Gene 49, 167-175, 1986
Gene 49, 167-175, 1986
A;Title: The human transferrin gene: 5' region contains conserved sequences which match
A;Reference number: A29090; MUID:87192006; PMID:3106157
A;Accession: A29090 A;Cross-references: UNIPROT:P02787; UNIPARC:UPI000002F089; EMBL:M12530; NID:g339452; PID:A;Note: the authors translated the codon CAA for residue 203 as Glu R;MacGillivray, R.T.A.; Mendez, E.; Shewale, J.G.; Sinha, S.K.; Lineback-Zins, J.; Brew, J. Biol. Chem. 258, 3543-353, 1983
A;Title: The primary structure of human serum transferrin. The structures of seven cyano A;Reference number: A92417; MUID:83160878; PMID:6833213
A;Accession: A92417 C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 15-Oct-1982 #sequence revision 30-Sep-1993 #text change 09-Jul-2004
C;Date: 15-Oct-1982 #sequence revision 30-Sep-1993 #text change 09-Jul-2004
C;Accession: A20981, A92417, A94044; A29090; A32739; I51959; I63133; I54011; I68160;
C;Accession: A2081, MCGill, J.R.; Moore, C.M.; Naylor, S.L.; van Bragt, P.H.; Bald Proc. Natl Acad. Sci. U.S.A. 81, 2752-2756, 1984
A;Title: Human transferrin: cDNA characterization and chromosomal localization.
A;Reference number: A20981; MUID:84194084; PMID:6585826
A;Contents: variant C A;Molecule type: mRNA A;Reeidues: 422-690,'G',692-698 <UZA> A;Cross-references: UNIPARC:UP1000016B0AB; EMBL:M12525; NID:g339468; PIDN:AAA61142.1; PI; R;MacGillivray, R.T.A.; Mendez, E.; Sinha, S.K.; Sutton, M.R.; Lineback-Zins, J.; Brew, I R;Park, I.; Schaeffer, B.; Sidoli, A.; Baralle, F.E.; Cohen, G.N.; Zakin, M.M. Proc. Natl. Acad. Sci. U.S.A. 82, 3149-3153, 1985
A;Title: Organization of the human transferrin gene: direct evidence that it originated A;Reference number: A94044; MUID:85216459; PMID:3858812 A; Accession: A94044 A; Note: the sequence shown A;Molecule type: protein A;Residues: 20-263,'E',265-328,'N',330-379,'SD',382-435,'D',437-557,'T',559-560,'P',562-A;Cross-references: UNIPARC:UPI0000174483 A; Molecule type: mRNA A; Residues: 1-698 < YAN> A; Accession: A20981 transferrin precursor [validated] N;Alternate names: siderophilin is the predominant electrophoretic genetic variant (C human or TfC)

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transferrin precursor - rabbit
(;Species: Oryctolegus cuniculus (domestic rabbit)
(;Species: Oryctolegus cuniculus (domestic rabbit)
(;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 22-Jun-1999
(;Accession: S16246; A61239; C61573; S00335; S02694; A26504; S14853
R;Banfield, D.K.; Chow, B.K.C.; Funk, W.D.; Robertson, K.A.; Umelas, T.M.; V
Blochim. Biophys. Acta 1089, 262-265, 1991
A;Title: The nucleotide sequence of rabbit liver transferrin cDNA.
A;Reference number: S16246; MUID:91274362; PMID:2054387
A;Accession: S16246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Map position: 3q21-3q21
A;Map position: 3q21-3q21
C;Function: 15/1; 72/3; 119/1; 168/1; 212/2; 231/1; 290/3; 350/1; 401/3; 433/1; 444/1; 49
C;Function: binds iron for delivery into cells
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron transport; metal binding; plasma
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-698/Product: transferrin #status experimental <MAT>
F;00-350/Domain: transferrin repeat homology <TRH1>
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A;Residues: 564-698 <RE4>
A;Residues: 564-698 <RE4>
A;Cross-references: UNIPARC:UPI000016B0AE; GB:M17614; NID:g339483; PIDN:AAA61148.1; PID:
C;Comment: Each of the two repetitive domains binds a ferric ion and a bicarbonate anion
C;Genetics:
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A;Molecule type: mRNA
A;Residues: 636-696 <RE2>
A;Residues: 636-696 <RE2>
A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:
A;Cross-references: UNPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:
R;Schaeffer, E.; Lucero, M.A.; Jeltsch, J.M.; Py, M.C.; Levin, M.J.; Chambon, P.; Cohen,
R;Schaeffer, E.; Lucero, M.A.; Jeltsch, J.M.; Py, M.C.; Levin, M.J.; Chambon, P.; Cohen,
Gene 56, 109-116, 1987
Gene 56, 109-116, 1987
A;Title: Complete structure of the human transferrin gene. Comparison with analogous chi
A;Reference number: I54011
A;Reference number: I54011
A;Reference number: I54011
A;Status: translated from GB/EMBL/DDBJ
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Proc. Natl. Acad. Sci. U.S.A. 86, 7260-7264, 1989
A;Title: Changes in brain gene expression shared by scrapie and Alzheimer disease.
A;Reference number: I48174; MUID:89386721; PMID:2780570
A;Accession: I63133
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A;Cross-references: GDB:120432; OMIM:190000
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A; Residues: 1-72 < RE3 >
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A;Molecule type: mRNA
A;Residues: 1-698 <RES>
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A;Title: A cloned gene for human transferrin.
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A;Title: The complete amino acid sequence of human serum transferrin.
A;Reference number: A93911, MUID:82222166; PMID:6953407
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A;Accession: I51959
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Best Local S
Matches 22
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-28-67,38-58,137-213,156-350,177-193,180-196,190-198,246-260,358-615,364-396,374-387,42
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95.7%;
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Pred. No. 2.8e-09;
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C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron transport; metal binding; plasma
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-694/Product: transferrin #status experimental <MAT>
F;19-349/Domain: transferrin repeat homology <TRH1>
F;355-682/Domain: transferrin repeat homology <TRH2>
F;27-66,37-57,136-212,155-349,176-192,179-195,189-197,245-259,357-611,363-395,373-386,420
F;508/Binding site: carbohydrate (Asn) (covalent) #status experimental
transferrin precursor - horse
N;Alternate names: growth-promoting factor
C;Species: Equus caballus (domestic horse)
C;Date: 06-Jan-1995 #sequence revision 06-Jan-1995 #text_change
C;Accession: S33761; S02145
R;Carpenter, M.A.; Broad, T.E.
Biochim. Biophys. Acta 1173, 230-232, 1993
A;Title: The CDNA sequence of horse transferrin.
A;Reference number: S33761; MUID:93277958; PMID:8504171
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A;Accession: S00335
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S33761
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A;Note: 516-Ile was also found
R;Heaphy, S.; Williams, J.
Biochem, J. 205, 611-617, 1982
A;Title: The preparation and partial characterization of N-terminal and C-terminal iron-t
A;Reference number: A26504; MUID:83074540; PMID:6816218
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FEBS Lett. 238, 39-42, 1988
A;Title: Evidence for a single glycan moiety in rabbit serum transferrin and location
A;Reference number: S02694; MUID:89005676; PMID:3169252
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A;Residues: 19-45,'S',47-48,'Y',50 <GOD>
A;Cross-references: UNIPARC:UPI00001744A8
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A;Residues: 19-26,'X',28-36,'X',38-53 <CHU>
A;Cross-references: UNIPARC:UPI00001744A7
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A; Residues: 19-24, 'N', 26, 'X', 28-29, 'S' <HEA>
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A; Residues: 482-515, 'V', 517-544 < EVA>
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A; Accession: C61573
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R;Chung, M.C.M.; Chan, S.L.; Shimizu, S.
Int. J. Biochem. 23, 609-616, 1991
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A;Cross-reference
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A; Title: Iron carrier proteins fa
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87.0%;
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Pred. No. 7.5e-08;
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; PMID:2065820
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C:Species: Mus musculus (house mouse)
C:Species: Mus musculus (house mouse)
C:Date: 30-Jun-1989 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C:Accession: A28446; A33482; B28438
R;Chen, L.H.; Bissell, M.J.
J. Biol. Chem. 262, 17247-17250, 1987
J. Biol. Chem. 262, 17247-17250, 1987
A;Title: Transferrin mRNA level in the mouse mammary gland is regulated by pregnancy and A;Reference number: A28446; MUID:88086992; PMID:3693348
A;Accession: A3482; MUID:90097932; PMID:2601714
A;Reference number: A33482; MUID:90097932; PMID:2601714
A;Accession: A33482; MUID:90097932; PMID:2601714
                                                                                                                                                                                                                     A;Molecule type: mRNA
A;Residues: 86-216 <PEN>
A;Cross-references: UNIPARC:UPI000016CEB1; GB:J03299; GB:J02737; NID:g198847; PIDN:AAA39
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication
                                                                                                                                                                                                                                                                                                                                                        R;Pentecost, B.T.; Teng, C.T.
J. Biol. Chem. 262, 10134-10139, 1987
A;Title: Lactotransferrin is the major estrogen inducible protein A;Reference number: A92596; MUID:87280033; PMID:3611056
A;Accession: B28438
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C;Keywords: duplication; glycoprotein; iron transport; metal binding; plasma
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-706/Product: transferrin #status experimental <WAT>
F;358-694/Domain: transferrin repeat homology <TRH2>
F;26-64,36-55,134-215,174-190,177-198,187-200,248-262,360-623,366-398,376-389,423-701,44
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A;Title: A growth-promoting factor for human myeloid leukemia A;Reference number: S02145; MUID:89076897; PMID:2909248
A;Accession: S02145
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A;Residues: 1-14;302-311 <IDZ>
A;Cross-references: UNIPARC:UPI0000177C57; UNIPARC:UPI0000177C58;
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A;Residues: 20-35,'X',37,'X',39-40,'X',43-44 <YOS:
A;Cross-references: UNIPARC:UPI0000177C56
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A;Molecule type: mRNA
A;Residues: 1-706 <CA
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                                              TTEDSIAKIMNGEADAMSLDGG
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                                                                                                      Score 92; DB 2;
Pred. No. 8e-07;
2; Mismatches
  216
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transferrin
C;Species: S
C;Date: 10-S
                                                                                            A;Note: sequence extracted from NCBI backbone (NCBIN:111151, NCBIP:111153)
R;Lydon, J.P.; O'Malley, B.R.; Saucedo, O.; Lee, T.; Headon, D.R.; Conneel:
Biochim. Biophys. Acta 1132, 97-99, 1992
A;Title: Nucleotide and primary amino acid sequence of porcine lactoferrin A;Reference number: S24173; MUID:92379101; PMID:1511016
A;Accession: S24173
                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Alexander, L.J.; Levine, W.B.; Teng, C.T.; Beattie, C.W. Anim. Genet. 23, 251-256, 1992
A;Title: Cloning and sequencing of the porcine lactoferrin cDNA A;Reference number: A45543; MUID:92367939; PMID:1503259
A;Accession: A45543
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A;Title: Isolation of transferrin from porcine gastric mucosa: comparison A;Reference number: A60520, MUID:90227903; PMID:2328566
A;Accession: A60520
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C;Date: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: S01384, A60520, Ā61573
R,Baldwin, G.S.; Weinstock, J.
Nucleic Acids Res. 16, 8720, 1988
A;Title: Nucleotide sequence of porcine liver transferrin.
A;Reference number: S01384; MUID:88335629; PMID:3419934
A;Accession: S01384
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A;Residues: 1-8,'X', 10-18,'XE', <CHU>
A;Cross-references: UNIPARC: UPI0000174486
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron transport;
F;1-696/Product: transferrin #status predicted <MAT>
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Int. J. Biochem. 23, 609-616, 1991
A;Title: Purification of transferrins and lactoferrin using DEAE Affi-Gel Blue A;Reference number: A61573; MUID:91293379; PMID:2065820
A;Accession: A61573
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A;Ross-references: UNIPROT:P09571; UNIPARC:UPI00001130C2; EMBL:X12386; NID:g2126; PIDN: A;Note: 308-Arg was also found
A;Molecule type: mRNA
A;Residues: 1-11,'W',13-50,'I',52-84,'G',86-120,'L',121-130,'I',132-282,'S',284-571,'Q',
A;Cross-references: UNIPARC:UPI000016C682; EMBL:M92089; NID:g164613; PIDN:AAA31102.1; PII
                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPROT:P14632; UNIPARC:UPI0000177C55
A;Experimental source: mammary gland
                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-703 < ALE>
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                                                                                                                                                                                                                                                                                                                                                                                                                  A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        lactoferrin precursor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 17-Feb-1994 #sequence_revision 17-Feb-1994 #text_change
C;Accession: A45543; S24173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Note: the authors suggest transferrin from gastric mucosa may R;Chung, M.C.M.; Chan, S.L.; Shimizu, S.
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A;Molecule type: mRNA
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Best Local
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Endocrinology 120, 332-340, 1987
A;Title: Transferrin messenger ribonucleic acid: molecular cloning and hormonal regulati
A;Reference number: A30014; MUID:87053639; PMID:3023031
A;Accession: A30014
A;Accession: A30014
A;Molecula type: mRNA
A;Residues: 518-687,'D',689-692,'TA',695 <HUG>
A;Cross-references: UNIPARC:UPI0000170B4C; GB:M27966; NID:g207439; PIDN:AAA42267.1; PID:
A;Title: The synthesis and secretion of rat transferrin.
A;Reference number: A14679; MUID:80049855; PMID:500689
A;Accession: A14679
A;Molecula type: protein
A;Residues: 20-47 <SCH>
A;Residues: 20-47 <SCH
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()Superiamily: transferrin; transferrin repeat homology
()Superiamily: transferrin; transferrin; iron binding; metalloprotein; milk
()Superiamily: transferrin; glycoprotein; iron binding; metalloprotein; milk
()F;1-19/Domain: signal sequence #status predicted <SIG>
()F;20-703/Product: lactoferrin repeat bomology <TRH1>
()F;20-350/Domain: transferrin repeat homology <TRH1>
()F;36-48/Region: antimicrobial
()F;36-48/Region: antimicrobial
()F;36-69/Domain: transferrin repeat homology <TRH2>
()F;36-62/B;362-312,171-187,184-195,245-259,362-394,372-385,419-698,439-661,471-546,49
()F;37,107,206,267/Binding site: iron (Asp. Tyr, Tyr, His) #status predicted
()F;470,447,540,609/Binding site: iron (Asp. Tyr, Tyr, His) #status predicted
()F;470,447,540,609/Binding site: carbonate (Arg) #status predicted
()F;490/Binding site: carbonate (Arg) #status predicted
()F;490/Binding site: carbonate (Asp. Tyr, Tyr, His) #status predicted
()F;490/Binding site: carbonate (Asp. Tyr, Tyr, His) #status predicted
A;Status: preliminary
A;Molecule type: protein
A;Residues: 89,'Y',91,'A',93-99,'V',101-102;'N',233,'AN',236-243;401-406,'N',408
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N;Alternate names: lung-derived growth factor; siderophilin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 16-Feb-1995 #sequence_revision 12-May-1995 #text_change 09-Jul-2004
C;Accession: S49163; S54980; A30014; A14679; A53289; A30512; I52203
R;Escrive, H.; Pierce, A.; Coddeville, B.; Gonzalez, F.; Benaissa, M.; Leger,
submitted to the EMBL Data Library, January 1994
A;Description: Rat mammary gland transferrin: glycan structure, nucleotide seq
A;Reference number: S49163
A;Accession: S49163
A;Status: preliminary
A;Status: preliminary
                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000177C51
R;Cavanaugh, P.G.; Nicolson, G.L.
J. Cell. Biochem. 47, 261-271, 1991
A;Title: Lung-derived growth factor that stimulates the A;Reference number: A53289; MUID:92165927; PMID:1791188
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A; Residues: 1-695 <ES2>
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A; Residues: 1-695 < ESC>
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Pred. No. 2e-06;
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A;Cross-references: UNIPARC:UPI00000EF28; UNIPARC:UPI00000E76D0; UNIPARC:UPI00000E7E7F
A;Experimental source: lung
A;Note: sequence modified after extraction from NCBI backbone
A;Note: sequence extracted from NCBI backbone (NCBIP:86115, NCBIP:86116, NCBIP:86114)
R;Purves, L., Linton, N.; Brandt, W.; Johnson, G.; Jacobs, P.
Biochim. Biophys. Acta 966, 318-327, 1988
A;Ticle: Properties of the transferrin associated with rat intestinal mucosa.
A;Reference number: A30512; MUID:88327006; PMID:3046665
A;Accession: A30512
A;Molecule type: protein
A;Residues: 20-30;639-643, KD',646, LKACD' <PUR>A;Cross-references: UNIPARC:UPI0000177C52; UNIPARC:UPI0000177C53
B;Aldred, A.R.; Howlett, G.J.; Schreiber, G.
Biochem. Biophys. Res. Commun. 122, 960-965, 1984
A;Title: Synthesis of rat transferrin in Escherichia coli containing a recombinant bacter A;Accession: I52203; MUID:84307580; PMID:6236811
A;Accession: I52203; MUID:84307580; PMID:6236811
                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: mRNA
A;Residues: 1-148, TT; 150-422, 'C', 424-711 <REY>
A;Residues: 1-148, TT; 150-422, 'C', 424-711 <REY>
A;Cross-references: UNIPARC:UPI000015C582; EMBL:X53961; NID:g34415; PIDN:CAA37914.1;
A;Cross-references: UNIPARC:UPI000015C582; EMBL:X53961; NID:g34415; PIDN:CAA37914.1;
R;Teng, C.T.; Liu, Y.; Yang, N.; Walmer, D.; Panella, T.

Mol. Endocrinol. 6, 1969-1981, 1992
Mol. Endocrinol. 6, 1969-1981, 1992
A;Title: Differential molecular mechanism of the estrogen action that regulates lacto
A;Reference number: A45401; MUID:93125571; PMID:1480183
A;Experimental source: placenta A;Note: sequence extracted from NCBI backbone (NCBIP:12 R;Powell, M.J.; Ogden, J.E. Nucleic Acids Res. 18, 4013, 1990 A;Title: Nucleotide sequence of human lactoferrin cDNA. A;Reference number: S10324; MUID:90326549; PMID:2374734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI000013737C; EMBL:U07643; NID:g467236; PIDN:AAB60324.1; R;Rey, M.W.; Woloshuk, S.L.; deboer, H.A.; Pieper, F.R. Nucleic Acids Res. 18, 5288, 1990

A;Title: Complete nucleotide sequence of human mammary gland lactoferrin.
A;Reference number: S11228; MUID:90384839; PMID:2402455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NyAlternate names: lactoferrin
C;Species: Homo sapiens (man)
C;Date: 31-Mar-1992 #sequence revision 21-Nov-1997 #text change 08-Dec-2000
C;Accession: G01394; S11228; Ā45401; S10324; S15853; S20841; S07160; A61169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-711 < CHO>
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A;Cross-references: UNIPARC:UPI00000E5C48, GB:M26113; NID:9207437; PIDN:AAA42266.1;
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A;Molecule type: mRNA
A;Residues: 7-25,'X',27-56,'A',58-64,267-295 <RES>
                                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-15 < TEN >
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                                                                                                                                                                                                                                  A;Cross-references: UNIPARC:UPI00000004DD; GB:S52659; NID:g263311;
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Pred. No.
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A;Note: this is the final paper in a series
R;Houen, G.; Hoegdall, E.V.; Barkholt, V.; Norskov, L.
Eur. J. Biochem. 241, 303-308, 1996
A;Title: Lactoferrin: similarity to diamine oxidase and pur
A;Reference number: S74119; MUID:97054624; PMID:8898921
A;Accession: S74119
A;Molecule type: protein
A;Residues: 'G',23-24,'R',26-27,'XX',30-32 <HOU>
A;Cross-references: UNIPARC:UPI000017448A
A;Experimental source: neutrophil granulocytes
C;Genetics:
C;Genetics:
C;Genetics: GB:LTF
                                                                                                                                                                                                                                                                           A;Cross-references: GDB:119368; OMIM:150210
A;Map position: 3q21-3q23
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron binding; milk
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-711/Product: lactotransferrin #status experimental <MAT>
F;21-356/Domain: transferrin repeat homology <TRH1>
F;21-356/Domain: transferrin repeat homology <TRH1>
F;360-699/Domain: transferrin repeat homology <TRH2>
F;350-699/Domain: transferrin repeat homology <TRH2>
F;350-699/Domain: cransferrin repeat homology <TRH2>
F;350-699/Domain: cransferrin repeat homology <TRH2>
F;350-699/Domain: decomposition repeat homology <TRH2>
F;350-699/Domain: cransferrin repeat homology <TRH2>
F;350-699/Domain: decomposition repeat homology <TRH2>
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A;Residues: 436-487, A', 489-711 <RAD>
A;Crose-references: UNIPARC:UPI000016ABE2; EMBL:M18642; NID:g186815; PIDN:AAA86665.1; R;Panella, T.J.; Liu, Y.; Eng, A.T.; Teng, C.T.
Cancer Res. 51, 3037-3043, 1991
A;Title: Polymorphism and altered methylation of the lactoferrin gene in normal leukoc A;Reference number: A61169; MUID:91235214; PMID:1674448
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R;Rado, T.A.; Wei, X.; Benz Jr., E.J.
Blood 70, 989-993, 1987
A;Title: Isolation of lactoferrin cDNA from
A;Reference number: S07160; MUID:88001031; F
A;Accession: S07160
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A;Residues: 20-140,142-169,171-203,'L',205,'K',207-208,'K',210-385,'Q',387-391,'W',393-A;Cross-references: UniPARC:UPI0000174489
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A;Residues: 3-711 <POW>
A;Cross-references: UNIPARC:UPI000016AC28; EMBL:X52941; NID:g34411; PIDN:CAA37116.1;
A;Cross-references: UNIPARC:UPI000016AC28; EMBL:X52941; NID:g34411; PIDN:CAA37116.1;
R;Stowell, K.M.; Rado, T.A.; Funk, W.D.; Tweedie, J.W.
Biochem. J. 276, 349-355, 1991
Biochem. J. 276, 349-355, 1991
A;Title: Expression of cloned human lactoferrin in baby-hamster kidney cells.
A;Reference number: S15853; MUID:91264786; PMID:2049066
A;Accession: S15853
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A;Residues: 20-28,'X',30-31 <ST2>
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A; Molecule type: mRNA
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A;Residues: 20-31 <ST1>
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                                                                                                                                             Local
                                                                                                                                                  Similarity
TTEDSIAKIMNGEADAMSLDGGF
                                                                                                                                             78.3%;
                                                                                                                                                  Score 90; DB 1;
Pred. No. 4.3e-06;
                                                                                                        Mismatches
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PMID:3477300
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carbonic anhydrase II inhibitor (transferrin homolog) precursor - pig C;Species: Sus scrofa domestica (domestic pig) C;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 09-Jul-2004 C;Accession: I47228
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N;Alternate names: lactotransferrin
C;Species: Mus musculus (house mouse
C;Date: 10-Sep-1999 #momnon
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E;1-19/Domain: signal sequence #status predicted <SIG-
E;20-707/Product: lactotransferrin #status predicted <MAT-
E;30-707/Domain: transferrin repeat homology <TRH2-
E;358-695/Domain: transferrin repeat homology <TRH2-
E;358-695/Domain: transferrin repeat homology <TRH2-
                                                                                                                                                      C;Keywords: duplication F;20-350/Domain: transferrin repeat homology <TRH1>
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A;Title: Characterization of estrogen-responsive mouse lactoferrin A;Reference number: A41205; MUID:92042099; PMID:1939212
A;Accession: A41205
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J. Biol. Chem. 262, 10134-10139, 1987
A;Title: Lactotransferrin is the major estrogen inducible protein of A;Reference number: A92596; MUID:87280033; PMID:3611056
A;Accession: A28438
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A;Title: Purification and characterization of a carbonic
A;Reference number: I47228; MUID:93099129; PMID:1463741
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C;Accession: A28438; A41205
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                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-704 < ROU>
                                                                                                                                                                                                                                                                                                                                                                A;Status: preliminary; translated from GB/EMBL/DDBJ
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Biochemistry 31, 12536-12542, 1992
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A; Residues: 1-15 <LIU>
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J. Biol. Chem. 2
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A; Residues: 3-707 < PEN>
                                                                                                                                                                                                    ;Superfamily: transferrin; transferrin repeat homology; Keywords: duplication
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                                                                                                                                   Query Match
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388
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                                                                                                          74.8%;
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Pred. No. 1.
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Pred. No. 8.9e-06;
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Bur. J. Biochem. 122, 291-295, 1982
A;Title: The complete nucleotide sequence of the chicken ovotransferrin mRNA.
A;Reference number: A91115; MUID:82138851; PMID:7060577
A;Accession: A91115
A;Acce
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A; Residues: 1-50,54-82,'V',84-92;102-146;162-168;170-228;241-283;289-333;338-344;350-351
A; Crestidues: 1-50,54-82,'V',04-92;102-146;162-168;170-228;241-283;289-333;338-344;350-351
A; Cross-references: UNIPARC:UPI0000118CCA; UNIPARC:UPI00001744AB; UNIPARC:UPI00001744B6; UNIPARC:UPI000001744B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC:UPI0000174B6; UNIPARC
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A;Title: The primary structure of hen ovotransferrin.
A;Reference number: A91116; MUID:82138852; PMID:6895872
A;Accession: A91116
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A;Residues: 1-23 <THI>
A;Cross-references: UNIPARC:UPI00001744AC
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A;Reference number: JC2323; MUID:94380047; PMID:8093048
A;Accession: JC2323
A;Molecule type: mRNA
A;Residues: 1-708 < LEP
A;Cross-references: UNIPARC:UPI0000177C50
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein
F;359-96/Domain: transferrin repeat homology < TRH2>
F;252,300,387,495,564/Binding site: carbohydrate (Asn) (
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C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 07-May-1999
C;Accession: JC2323
R;Le Provost, F.; Nocart, M.; Guerin, G.; Martin, P.
Biochem. Biophys. Res. Commun. 203, 1324-1332, 1994
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Pred. No. 3.9e-0
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C;Comment: C
C;Comment: I
C;Comment: I
C;Genetics:
                C;Species: Bos primigenius taurus (cattle)
C;Date: 31-Mar-1992 #sequence_revision 21-Nov-1997 #text_change 09-Ju
C;Accession: I45919; S14674; S14110; S18517; JT0595; S13097; S18518;
R;Tsang, T.C.; Burns, D.K.; Wang, F.; Pan, Y.
FASEB J. 6, 233, 1991
A;Title: Cloning of a 80-kD advanced glycosylation end product (AGE)
A;Reference number: I45919
A;Accession: I45919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; egg white; glycoprotein; iron binding; plasma
F;1-19/Domain: signal sequence #status experimental <SIC>
F;20-705/Product: transferrin #status experimental <MAT>
F;21-35/Domain: transferrin repeat homology <TR11>
F;35-502/Domain: transferrin repeat homology <TR11>
F;35-692/Domain: transferrin repeat homology <TR12-
F;35-692/Domain: transferrin repeat homology <TR16-
F;39-63,134-216,179-193,190-201,247-261,367-399,424-699,440-662,473-549,497-690/Disulfide
F;492/Binding site: carbohydrate (Asn) (Covalent) (partial) #status experimental
F;492/Binding site: carbohydrate (Asn) (Covalent) (spartial) #status predicted
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A;Residues: 20-23:295-302:336-366;674-679,'T',681,'F',683-685,'N',687-705 <WIZ>A;Residues: 20-23:295-302:336-366;674-679,'T',681,'F',683-685,'N',687-705 <WIZ>A;Cross-references: UNIPARC:UPI00001744BE; UNIPARC:UPI00001744BF; UNIPARC:UPI00001744CO; C;Comment: Ovotransferrin (conalbumin) and transferrin have identical protein components rrin is synthesized in the liver.
C;Comment: Synthesized in the liver.
C;Comment: Dlasma apotransferrin promotes the oxidation of ferrous ions, which would othe C;Comment: In electrophoretic and genetic studies, transferrin shows strong polymorphism,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biochem. J. 116, 515-532, 1970
A,Title: The amino acid sequences of cysteic acid-containing peptides from A,Reference number: A90246; MUID:70141846; PMID:4907959
A,Contents: annotation; disulfide bonds
R;Williams, J.; Moreton, K.
Biochem. J. 251, 849-855, 1988
Biochem. J. 251, 849-855, 1988
A,Title: The dimerization of half-molecule fragments of transferrin.
A,Reference number: S02476; MUID:88326225; PMID:3415649
A,Accession: S02476
                                                                                                                                                                                                                        N;Alternate names:
C;Species: Bos prin
C;Date: 31-Mar-1992
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A;Title: The amino acid sequence of a carbohydrate-containing
A;Reference number: A90282; MUID:76039467; PMID:1172663
A;Accession: A90282
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A; Residues: 20-28, 'X', 30-38, 'X', 40-43, 'S'
A; Cross-references: UNIPARC: UPI00001744BC
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Int. J. Biochem. 23, 609-616, 1991
A;Title: Purification of transferrins and lactoferrin using
A;Reference number: A61573; MUID:91293379; PMID:2065820
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A; Residues: 20-28, 'X', 30-38, 'X', 40-44 <GENS
A; Cross-references: UNIPARC: UDI00001744BB
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A;Status: translated from GB/EMBL/DDBJ
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8518; S13881;
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A; Molecule type: mRNA
A; Residues: 1-65, 'PG', 68-296, 'S', 298-339, 'A', 341-708 <GOO>
A; Cross-references: UNIPARC: UPI0000174495; GB:M63502
A; Note: the authors translated the codon CCG for residue 66 at R; Mead, P.E.; Tweedie, J.W.
Nucleic Acids Res. 18, 7167, 1990
A; Title: cDNA and protein sequence of bovine lactoferrin.
A; Reference number: S13097; MUID:91088328; PMID:2263492
A; Accession: S13097
A; Molecule type: mRNA
A; Residues: 28-33, 'DS', 36-38, 'P', 40-708 <MEA>
A; Cross-references: UNIPARC: UPI0000174496; EMBL:X54801
A;Cross-references: UNIPARC:UPI00000475A8
R;Shimazaki, K.; Tanaka, T.; Kon, H.; Oota, K.; Kawaguchi,
J. Dairy Sci. 76, 946-955, 193
A;Title: Separation and characterization of the C-terminal
A;Reference number: A56659; MUID:93253156; PMID:8486845
                                                                                                                                                          A;Molecule type: protein
A;Residues: 20-27'X',29-37,'X',39-54,'X',56-59 <REJ>
A;Cross-references: UNIPARC:UPI00001744A0
R;Bellamy, W.; Takase, M.; Yamauchi, K.; Wakabayashi, H.; Kawase, Biochim. Biophys. Acta 1121, 130-136, 1992
A;Title: Identification of the bactericidal domain of lactoferrin.
A;Reference number: S21756; MUID:92287941; PMID:159934
A;Accession: S21756
A;Molecule type: protein
A;Residues: 36-60 <BEL>
A;Residues: 36-60 <BEL>
A;Molecule type: Protein
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Comp. Biochem. Physiol. B 93, 929-934, 1989
A;Title: Purification and characterization of bovine lactoferrin
A;Reference number: PL0148; MUID:90031466; PMID:2805645
A;Accession: PL0148
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A;Residues: 20-47;59-66;132-139;256-277;278,305-332;343-351;361-363;586,587-589;598-619
A;Cross-references: UNIPARC:UPI0000174493; UNIPARC:UPI0000174497; UNIPARC:UPI0000174498
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A;Residues: 20-35;82-114;148-163,'PP',166-178,'V','P';183-190;205-212;230-239;304-339;59
A;Cross-references: UNIPARC:UPI000017448C; UNIPARC:UPI000017448D; UNIPARC:UPI000017448E;
493; UNIPARC:UPI0000174494
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A; Residues: 28-38,'P', 40-86,'C', 88-708 <ME3>
A; Cross-references: UNIPARC: UPI000017449F; EMBL: X54801
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A;Residues: 3-144,'V',146-339,'A',341-438,'Y',440-513,'R',515-708
A;Cross-references: UNIPARC:UPI000017448B; EMBL:X57084
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A; Residues: 1-708 < TS
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;Cross-references: UNIPARC:UPD1000016C335; EMBL:X57084; NID:9505; PIDN:CAA40366.1;
;Pierce, A.; Colavizza, D.; Benaissa, M.; Maes, P.; Tartar, A.; Montreuil, J.; Sp
ur. J. Biochem. 196, 177-184, 1991
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Biophys. Res. Commun. 180, 75-84, 1991
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F;252,300,307,307,
F;414,452,545,614/Binding site: 1100.
F;482/Binding site: carbonate (Arg) #status
                                                                                                                                                                                                                                                      C;Species: Salmo sal
C;Date: 16-Jul-1999
C;Accession: T11749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
S12100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 20-25;302-308;359-366,'X',368-376,'X',378 <SHI>
A;Residues: 20-25;302-308;359-366,'X',368-376,'X',378 <SHI>
A;Cross-references: UNIPARC:UPI00001744A1; UNIPARC:UPI00001744A2; UNIPARC:UPI000
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication; glycoprotein; iron; iron binding; metalloprotein; milk
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-708/Product: lactotransferrin #status experimental <MAT>
                                                                                                                                       R;Kvingedal, A.M.; Rorvik, K.A.; Alestrom, P.
Mol. Marine Biol. Biotechnol. 2, 233-238, 1993
A;Title: Cloning and characterization of Atlantic salmon
A;Reference number: 217332; MUID:94122797; PMID:8293074
A;Accession: T11749
                                                                                                                                                                                                                                                                                                                                                           RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: S12100
R;Moskaitis, J.E.; Pastori, R.L.; Schoenberg, D.R.
Nucleic Acids Res. 18, 6135, 1990
A;Title: The nucleotide sequence of Xenopus laevis transferrin mRNA
A;Reference number: S12100; MUID:91045087; PMID:2235503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Xenopus laevis (African clawed frog)
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995
C;Accession: S12100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F;38-55/Disulfide bonds: #status predicted
F;79,111,211,272/Binding site: iron (Asp, Tyr, Tyr, His) #status e
F;140/Binding site: carbonate (Arg) #status experimental
F;252,300,387,495,564/Binding site: carbohydrate (Asn) (covalent)
F;414,452,545,614/Binding site: iron (Asp, Tyr, Tyr, His) #status
                                                                    A; Molecule type: mRNA
A; Residues: 1-690 < KVI>
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transferrin precursor - African clawed frog)
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                        A; Experimental source: liver
                                            A; Cross-references: UNIPROT: P80426;
                                                                                                                 A; Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-717 < MOS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: S12100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;359-696/Domain: transferrin repeat homology <TRH2>;28-64,134-217,176-192,179-200,189-202,250-264,367-399,377-390,424-703,444-666,476-551
;Superfamily: transferrin;
                                                                                                                                                                                                                                                                                   Date: 16-Jul-1999 #sequence
                                                                                                                                                                                                                                                                                                            ransferrin - Atlantic salmon
;Species: Salmo salar (Atlantic salmon)
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56.5%;
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Pred. No. 0.0005;
6; Mismatches
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                                              UNIPARC: UPI0000137362;
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115
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Q7YS20-PIG
Q8MNNB-PIG
TRFE RAT
Q7TWX0 RAT
Q7TWX0 RAT
Q7TWC7-RAT
Q7TWC7-RAT
Q7TPC4-RAT
Q7TPC4-RAT
Q7TPC4-MOUSE
Q3TP24-MOUSE
Q4FUR3-MOUSE
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Q7tsx8 marmota mon
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P27425 equus cabal
P09571 sus scrofa
Q921i1 mus musculu
Q3ubw7 mus musculu
Q58e69 m transfer
P14632 sus scrofa
Q6yt39 sus scrofa
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Q7ys20 sus scrofa
Q8wmn8 sus scrofa
Q12346 rattus norv
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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

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Rieder M.J., Carrington D.P., da Ponte S.
Ahearn M.O., Kuldanek S.A., Rajkumar N.,
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Fairbanks V.F.;
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Williams P., Dehoff B., Dunn P., O'Neal K.L.,
"A cloned gene for human transferrin.";
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MEDLINE=88056305; Pu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96:4071-4074 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            [LARGE SCALE MRNA]
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L., Riemen M.W., Tice P.
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n E.J., Yi Q.,
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seven cyanogen bromide fragments and t structure.";
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MEDLINE=87192006; PubMed=3106157; DOI=10.1016/0378-1119(86)90277-5; Adrian G.S., Korinek B.W., Bowman B.H., Yang F.;

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                                                                                                                                                                      "Alternative splicing prevents transferrin differentiation of a human oligodendrocyte J. Neurosci. Res. 61:388-395(2000).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        originated
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Lineback-Zins J., Brew K.;
"The primary structure of human
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MEDLINE=84153910; PubMed=6322780;
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Proc. Natl. Acad. Sci. U.S.A.
                                                                                                                                    TISSUE=Brai
                                                                                                                                                                                                                                                               NUCLEOTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Organization of the human transferrin originated by gene_duplication.";
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Yu Y., Zh
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216459; PubMed=3858812;
chaeffer E., Sidoli A.,
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24-MAY-2005,
                            24-MAY-2005, sequence version 1.
07-FEB-2006, entry version 4.
Transferrin variant (Fragment).
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MEDLINE=98434369; PubMed=9760232; DOI=10.1021/bi9812064;
MEDLINE=98494369; PubMed=9760232; DOI=10.1021/bi9812064;
Moodworth R.C., Baker E.N.;
"Ligand-induced conformational change in transferrins: crystal
                                                                                                                                                                                                                                                                                                                                                                                                                                        X-RAY CRYSTALLOGRAPHY (1.6 ANGSTROMS) OF 22-350.

MEDLINE=98272665; PubMed=9609685; DOI=10.1021/bi980355j;

MAGGIllivray R.T.A., Moore S.A., Chen J., Anderson B.F., Ba
Luo Y., Bewley M.C., Smith C.A., Murphy M.E.P., Wang Y., Ma
Woodworth R.C., Brayer G.D., Baker E.N.;

"Two high-resolution crystal structures of the recombinant
human transferrin reveal a structural change implicated in
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Moore R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=92031536; PubMed=1932003; Woodworth R.C., Mason A.B., Funk W.D., McGillivray R.T.A.; Woodworth R.C., Mason A.B., Eunk W.D., McGillivray R.T.A.; Expression and initial characterization of five site-directed of the N-terminal half-molecule of human transferrin.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=96007936; PubMed=7498159; Kovalyov L.I., Shishkin S.S., Efimochkin A Ershova E.S., Egorov T.A., Musalyamov A.K. "The major protein expression profile and database of human heart.";
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Duguid J.R., Bohmont C.W., Liu N.G., Tourtellotte W.
"Changes in brain gene expression shared by scrapie
disease.";
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J. Proteome Res. 4:2070-2080(2005).
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MEDLINE=82222166; PubMed=6953407;
   Eukaryota;
                                                                                                                                                                                                                                                                                                          structure of
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                 Homo sapiens
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Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                             integrated into UniProtKB/TrEMBL
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                    (Human)
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SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 2; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
NON TER 1 1
SEQUENCE 698 AA; 77080 MW; A541
                                                                                                                  Q4R5M1;
19-JUL-2005, integrated into UniProtKB/TrEMBL.
19-JUL-2005, sequence version 1.
19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 5.
Brain cDNA, clone: QccE-13766, similar to human transferrin (TF), .
Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
Cercopithecidae; Cercopithecinae; Macaca.
NUCLEOTIDE SEQUENCE.
PubMed=15944441; DOI=10.1093/molbev/msi187;
Osada N., Hirata M., Tanuma R., Kusuda J.,
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GO; GO:000819; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
PANYHER; PTHR11485; Peptidase S60;
PANYHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 2.
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                                                                                                      NCBI_TaxID=9541;
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Best Local S
Matches 22
HSSP; P19134; JUNF:
HSSP; P19134; JUNF:
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; F:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60,
PANTHER; PTHR11485; Peptidase_S60; 1.
Pfam; PF00405; Transferrin; 2.
PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR, FER; 2.
PROSITE; PS000205; TRANSFERRIN_1; 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rinaudo J.A.S., Gerin J.L.; Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9995;
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SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN_1; 2.

PROSITE; PS00205; TRANSFERRIN_2; 2.

PROSITE; PS00207; TRANSFERRIN_3; 2.

PROSITE; PS00207; TRANSFERRIN_3; 2.
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Submitted (JUN-2005) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
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GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
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GO:0005576; C:extracellular region;
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95.7%;
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Pred. No. 7.
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA.
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Attribution-NoDerivs License
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Best Local
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p19134; 046514;
01-NOV-1990, integrated into Un
15-JUL-1999, sequence version 4
07-MAR-2006, entry version 61.
Serotransferrin precursor (Tran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE;
PROSITE;
SEQUENCE
 preliminary structure analysis of the N-terminal half A resolution.";
                                                                                      MEDLINE=89026775; PubMed=3179277;
Bailey S., Evans R.W., Garratt R.C., Gorinsky B., Hasnain S.,
Horsburgh C., Jhoti H., Lindley P.F., Mydin A., Sarra R., Wate
"Molecular structure of serum transferrin at 3.3-A resolution.
Biochemistry 27:5804-5812(1988).
                                                                                                                                                                                         "Evidence for a single glycan moiety in rabbit serum transferrin location of the glycan within the polypeptide chain."; FEBS Lett. 238:39-42(1988).
                                                                                                                                                                                                                              Evans R.W.,
                                                                                                                                                                                                                                                                                                                    MEDLINE=88209278; PubM
Godovac-Zimmermann J.;
                                                                                                                                                                                                                                                                                                                                                                                   encoding gene."
                                                                                                                                                                                                                                                                                                                                                                                               "Cloning and structural organisation
                                                                                                                                                                                                                                                                                                                                                                                                                     Ghareeb B.A.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biochim. Biophys. Acta 1089:262-265(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=91274362; PubMed=2054387; DOI=10.1016/0167-4781(91)90021-D; Banfield D.K., Chow B.K.-C., Funk W.D., Robertson K.A., Umelas T.M., Woodworth R.C., Macgillivray R.T.A.; "The nucleotide sequence of rabbit liver transferrin cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryctolagus
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     binding globulin)
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                                                                                                                                                     TISSUE=P
                                                                                                                                                                 X-RAY CRYSTALLOGRAPHY (3.3 ANGSTROMS).
                                                                                                                                                                                                                                         MEDLINE=89005676; PubMed=3169252; DOI=10.1016/0014-5793(88)80221-7;
                                                                                                                                                                                                                                                                                                          "Isolation, characterization and N-terminal amino-acid
                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=New
                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Liver
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                                                                                                                                                                                                                                                       PROTEIN SEQUENCE OF 483-545
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Crystallogr.
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                                                                                                                                                                                                                                                                                           transferrin."
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PS00207; TRANSFERRIN_3;
694 AA; 76466 MW; 40
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                                                                                                                                                       lasma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STEDCIAKIMNGEADAMSLDGGY 414
                                                                                                                                                                                                                                                                                                                                                                        (OCT-1997) to the
                                                                                                                                                                                                                                                                                                                                                                                                                     Zealand white;
Thepot D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE [GENOMIC DNA].
                                                                                                                                                                                                                                Aitken A., Patel K.J
                                                                                                                                                                                                                                                                                 Hoppe-Seyler 369:93-96(1988)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    precursor (Transferrin) (Siderophilin)
                                                                                                                                                                                                                                                                                                                                   PubMed=3365331;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 103;
Pred. No. 8.
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3; 2.
40053F7DC1CFCA87 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                of the rabbit transferrin
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                          N-terminal half-molecule
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DR EMBL; AF03161; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031611; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031613; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031613; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031615; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031615; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031616; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031619; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031619; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031620; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031621; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031621; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031622; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031623; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031624; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031625; AAB94136.1; JOINED; Genomic DNA.

DR EMBL; AF031627; TRAASPERTIN 3:

DR MERODS; S60.972; --

DR PANTHER; PTHR11485; Peptidase S60; 1.

DR PANTHER; PTHR11485; Peptidase S60; 1.

DR PANTHER; PFO0405; Transferrin; 2.

DR PANTHER; PRO0405; Transferrin; 2.

DR PARTS; SM00094; TR FER; 2.

DR PROSITE; PS00206; TRANSFERRIN 3: 2.

DR PROSITE; PS00207; TRANSFERRIN 3: 2.

DR PROSITE; PS002
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DOMAIN
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-i- SUBUNIT: Monomer.
-i- SUBCELLULAR LOCATION: Secreted protein.
-i- TISSUE SPECIFICITY: Expressed by the liver and secreted in plasma.
-i- MISCELLANEOUS: The peptidase S60 domains do not have protease function as the catalytic residues are missing.
-i- SIMILARITY: Belongs to the transferrin family.
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                                          Serotransferrin.
/FTId=PRO_000035717.
Peptidase 560 1.
Peptidase 560 2.
Iron 1.
Iron 1.
Iron 1.
Iron 2.
Iron 3.
Iron 4.
Iron 4
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Matches 20; Conserv
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Missing (in Ref. 1).
K -> S (in Ref. 3).
P -> Y (in Ref. 3).
      Score 101; DB 1;
Pred. No. 1.8e-06;
1; Mismatches 2
                                          Length 695;
        Indels
        0
      Gaps
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0;

TTEDSIAKIMNGEADAMSLDGGF

23

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PIGE CONSTRUCTION OF THE PROPERTY OF A REAL PROPERTY OF THE PR
                                                                                                                                                                                                                            PANTHER; FILE.

PEAM; PF00405; Transfer.....

PINTS; PR00422; TRANSFERRIN.

SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN 2; 2.

PROSITE; PS00206; TRANSFERRIN 3; 2.

PROSITE; PS00207; TRANSFERRIN 3; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Equus caballus (Horse).
Eukaryota; Metazoa; Cho:
Mammalia; Eutheria; Lau
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HORSE
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P27425;
01-AUG-1992, integrated into U
01-AUG-1992, sequence version
21-FEB-2006, entry version 48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001156; Peptidase_S60. PANTHER; PTHR11485; Peptidase_S60; Pfam; PF00405; Transferrin; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; M69020; AAA30958.1; -; mRNA.
EMBL; U21127; AAA63684.1; -; mRNA.
PIR; S33761; S33761.
HSSP; P02787; LA8E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=93277958; PubMede=8504171; DOI=10.1016/0167-4781(93)90186-H; Carpenter M.A., Broad T.E.;
"The cDNA sequence of horse transferrin."; Biochim. Biophys. Acta 1173:230-232(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Serotransferrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       McDowell K.J., Adams M.H., Baker C.B.;
Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    binding globulin).
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SUBCELULAR LOCATION: Secreted protein.

TISSUE SPECIFICITY: Expressed by the liver and a miscellaneous: The peptidase S60 domains do not function as the catalytic residues are missing.

SIMILARITY: Belongs to the transferrin family.

SIMILARITY: Contains 2 peptidase S60 domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION: Transferrins are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate. It is responsible for the transport of iron from sites of absorption and heme degradation to those of storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.
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Chordata; Craniata; Vertebrata;
Laurasiatheria; Perissodactyla;
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Iron 1
Iron 1
Iron 1
Iron 2
Iron 2
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Peptidase S60 1.
Peptidase S60 2.
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(By similarity).
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                                      394
                                                                            Similarity
                                      STEDCIAKIVKGEADAMSLDGGF 416
                                                    TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                          508
519
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                                                                    Conservative
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                                                                            83.5%;
                                                                                                   MW;
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By similarity.
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                                                                            Score 96;
Pred. No.
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By similarity.
By similarity.
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By similarity.
1A0FA566C0409DBA CRC64;
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Similarity).
Carbonate 2; via amide nitrogen
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Carbonate 2
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PRT;
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ches 2;
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similarity)
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TRRE PIG STANDARD,

P09571;

01-JUL-1989, integrated into Unif

01-FEB-1991, sequence version 2.

21-FEB-2006, entry version 59.
PROTEIN SEQUENCE OF 1-20.

MEDLINE=9129379; PubMed=2065820;

Chung M.C., Chan S.L., Shimizu S.;

"Purification of transferrins and lactoferrin using DEAE affi-gel
                                                               MEDLINE-88335629; PubMed-3419934;
Baldwin G.S., Weinstock J.;
"Nucleotide sequence of porcine liver transferrin.";
Nucleic Acids Res. 16:8720-8720(1988).
                                                                                                                                                                             Eukaryota;
Mammalia;
                                                                                                                                                                                                      Sus scrofa
                                                                                                                                                                                                                               globulin).
                                                                                                                  TISSUE=Liver;
                                                                                                                              NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                   Name=TF;
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a; Metazoa;
; Eutheria;
                                                                                                                                                                                                                                                                                integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                             Chordata; Craniata; Vertebrata; Euteleostomi; Laurasiatheria; Cetartiodactyla; Suina; Suidae;
                                                                                                                              [MRNA]
                                                                                                                                                                                                                                           (Siderophilin)
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                                                                                                                                                                                                                                            (Beta-1
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PROTEIN SEQUENCE OF 1-15.

MEDLINE=90227903; PubMed=2328566;
Baldwin G.S., Bacic T., Chandler R., Grego B.,
Simpson R.J., Toh B.H., Weinstock J.;

"Isolation of transferrin from porcine gastric porcine serum transferrin.";

Comp. Biochem. Physiol. 95B:261-268(1990).
[4]
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PDB; 1H76; X-ray; A=1-696.
InterPro, IPR001156; Peptidase_S60; 1.
PANTHER; PTHR11485; Peptidase_S60; 1.
Pfan; PF00405; Transferrin; 2.
Pfan; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN_1; 2.
PROSITE; PS00206; TRANSFERRIN_2; 2.
PROSITE; PS00206; TRANSFERRIN_3; 2.
D-Structure; Direct protein sequencing; Glycoprotein; Ion transport; Metal-binding; Methylation; Polymorphism;
                                                          BINDING
BINDING
                                                                                                                                                                     DOMAIN
METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -I- FUNCTION: Transferring are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate. It is responsible for the transport of iron from sites of absorption and heme degradation to those of storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.

-I- SUBCELLULAR LOCATION: Secreted protein.
-I- TISSUE SPECIFICITY: Expressed by the liver and secreted in plasma.
-I- MISCELLANEOUS: The peptidase S60 domains do not have protease function as the catalytic residues are missing.
-I- SIMILARITY: Belongs to the transferrin family.
-I- SIMILARITY: Contains 2 peptidase S60 domains.
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Hall D.R., Hadden J.M., Leonard G.A.,
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Crystallogr. D 58:70-80(2002)
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Peptidase S60 2.
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STRAIN-C57BL/6J; TISSUE-Mammary gland;
PubMed=16141072; DOI=10.1126/science.1112014;
Carninci P., Kasukawa T., Katayama S., Gough J., Frith M.C., Maeda l
Oyama R., Ravasi T., Lenhard B., Wells C., Kodzius R., Shimokawa K.
Bajic V.B., Berenner S.E., Batalov S., Forrest A.R., Zavolan M.,
Davis M.J., Wilming L.G., Aidinis V., Brenst R.N., Bailey T.L.,
Ambesi-Impiombato A., Apweiler R., Aturaliya R.N., Bailey T.L.,
Bansal M., Baxter L., Beisel K.W., Bersano T., Bono H., Chalk A.M.,
Chiu K.P., Choudhary V., Christoffels A., Clutterbuck D.R.,
                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                     Serotransferrin precure binding globulin).
Name=Tf; Synonyms=Trf;
                                                                                                                                                                                                                                                                                                             06-JUN 2002, integrated into UniProtKB/Swiss-Prot. 01-DEC-2001, sequence version 1. 07-MAR-2006, entry version 44. Serotransferrin precursor (Transferrin) (Siderophi
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                                                                                                                                                                                        STRAIN-BALB/c;
                                                                                                                                                                                                                        NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
                                                                                                                NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
                                                                                                                                                                               Lai D.-Z.;
                                                                                                                                                                                                       NUCLEOTIDE
                                                                                                                                                               Construction of a robust CHO cell-line for biopharmaceutical
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                                                                                                                                                                                        SEQUENCE.
B/c; TISSUE=Liver;
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RA Crowe M.L., Dalla E., Dalrymple B.P., de Bono B., Della Gatta G., RA di Bernardo D., Down T., Engstrom P., Fagiolini M., Faulkner G., RA Fletcher C.F., Fukushima T., Furuno M., Futaki S., Gariboldi M., Rad Georgii-Hemming P., Gingeras T.R., Gojobori T. Green R.E., RA Hill D., Huminiecki L., Iacono M., Ikeo K., Iwama A., Ishikawa T., RA Hill D., Huminiecki L., Iacono M., Ikeo K., Iwama A., Ishikawa T., RA Jakt M., Kanapin A., Katch M., Kawasawa Y., Kelso J., Kitahura H., Kurochkin I.V., Lareau L.F., Lazarevic D., Lipovich L., Liu J., RA Kurochkin I.V., Lareau L.F., Lazarevic D., Lipovich L., Liu J., RA Kurochkin I.V., Lareau L.F., Lazarevic D., Lipovich L., Liu J., RA Matsuda H., Matsuzawa S., Miki H., Mignone F., Miyake S., Morris K., RA Milsson R., Nishiguchi S., Nishikawa S., Nori F., Ohara O., Olazado V., Pang K.C., Pavan W.J., Pavesi G., Pesole G., Patrovsky N., Piazza S., Reed J., Reid J.F., Ring B.Z., Ringwald M., Patrovsky N., Piazza S., Reed J., Reid J.F., Ring B.Z., Ringwald M., Schneider C., Schiguchi K., Semple C.A., Seno S., Seesa L., Sheng Y., RA Rost B., Ruan Y., Salzberg S.L., Sandelin A., Schneider C., Schipachi K., Jandelin R., Sultana R., Takenaka Y., Taki K., RA Tammoja K., Tan S.L., Tang S., Taylor M.S., Tegner J., Teichmann S.A., Wahlestedt C., Mattick J.S., Hume D.A., Kai C., Sasaki D., Tomaru Y., RA Wahlestedt C., Mattick J.S., Hume D.A., Kai C., Sasaki D., Tomaru Y., RA Tida J., Tammuza K., Itoh M., Kato T., Kawaji H., Kawagashira N., RA Taylon T., Kojima M., Kondo S., Konno H., Nakano K., Ninomiya N., RA Havashizaki Y., Chada M., Plessy C., Shibata K., Shiraki T., Suzuki H., Kawai J., Ra Havashizaki Y., Stathiki A., Okamura-Oho Y., Suzuki H., Kawai J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                               RX MEDLINE=2238257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altachul S.F., Zeeberg B., Buerow K.H., Schaefer C.F., Bhat N.K.,
RA Altachul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human rand mouse cDNA semences."

"Generation and initial analysis of more than 15,000 full-length human rand mouse cDNA semences."
pregnancy and J. Biol. Chem.
                                                                                                                                                                                                                Chaudhary J., Skinner M.K.; "Comparative sequence analysis promoters: hormonal regulation
                                                                                                                                                                                                                                                                                   MEDLINE=98284323; PubMed=9621303;
DOI=10.1002/(SICI)1098-2795(199807)50:3<273::AID-MRD3>3.3.CO;2-C;
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                                                                      Chen L.-H., Bissell M.J.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     and mouse cDNA sequences.";
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                                          "Transferrin mRNA level
                                                                                          MEDLINE=88086992;
                                                                                                                     NUCLEOTIDE SEQUENCE
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mRNA level in the mouse extracellular matrix."; 262:17247-17250(1987).
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                                                                                               PubMed=3693348;
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Ensembl; ENSWUSGO0000032554; Mus musculus.
MGI; MGI:9821; Trf.
GO; GO:0030139; C:endocytic vesicle; IDA.
GO; GO:0005768; C:endosome; IDA.
GO; GO:0005615; C:extracellular space; TAS.
InterPro; IPRO1156; Peptidase S60;
PANTHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; BC008559; AAH08559.1; -; mRNA.
EMBL; AF027336; AAB84034.1; -; Genomic
EMBL; M23014; AAA40489.1; -; mRNA.
EMBL; M23015; AAA40490.1; -; mRNA.
EMBL; M23016; AAA40491.1; -; mRNA.
EMBL; M23016; AAA40491.1; -; mRNA.
EMBL; S67217; AAB28966.2; -; mRNA.
EMBL; J03299; AAA39438.1; -; mRNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted
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MEDLINE=87280033; PubMed=3611056;
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MEDLINE=94068311;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBUNIT: Monomer.
SUBCELLULAR LOCATION: Secreted protein.
TISSUE SPECIFICITY: Expressed by the liver and INTISSUE State of the transferrin family.
SIMILARITY: Contains 2 peptidase S60 domains.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.
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AK085754, BAC39532.1; -; mRNA.
BC012313, AAH12313.1; -; mRNA.
BC022986; AAH22986.1; -; mRNA.
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/FTId=PRO_0000035716.
Peptidase S60 1.
Peptidase S60 2.
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Iron 1 (By similarity).
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Matches 18
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DISULFID
                                                                                                                                                                                                                                                                                                              STRAIN-C57BL/6J; TISSUE-Bone marrow; PubMed=16141072; DOI=10.1126/science. Il112014; PubMed=16141072; DOI=10.1126/science. Gondh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Muroidea; Muridae; Murinae; Mus.
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MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
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, Kitamura H.,
merfeld S.K.,
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RX MEDLINE=22154683; PISSUE=BORE MAIROW;
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Schriml L.M., Kanapin A., Mateuda H., Batalov S., Beisel K.W.,
RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA Ravasi T., Reed D.G., Reed D.G., Reid J., Ring B.Z., Ringwald M.,
RA Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
RA Wilming L.G., Wynshaw-Boris A., Yangisawa M., Yang I., Yang I.,
RA Wilming L.G., Wynshaw-Boris A., Yangisawa M., Yang I., Yang I.,
RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA Miyazaki A., Sakai K., Kawai J., Aizawa K., Arakawa T., Fukuda S.,
RA Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,
RA Miyazaki A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Miyazaki A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Mariya A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Mariya A., Yang L., Carninci P., Hayatsu N.,
RA Waterston R., Shinagawa A.,
RA Waterston R., Waterston R., Lander E.S., Rogers J.,
RA Mariya A., Yang L., Rad M., Waterston R., Lander E.S., Rogers J.,
RA Mariya A., Yang L., Rad M., Waterston R.
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Liuni S., McWilliam S., Madan Babu M., Madora M., Marchionni L.,
An Matsuda H., Matsuzawa S., Miki H., Mignone F., Miyake S., Morris K.,
An Mottagui-Tabbar S., Mulder N., Nakano N., Nakauchi H., Ng P.,
An Nilsson R., Nishiguchi S., Nishikawa S., Nori F., Ohara O.,
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An Shibata Y., Shimada H., Shimada K., Silva D., Sinclair B.,
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An Vamanishi H., Zabarovsky E., Zhu S., Zimmer A., Hide W., Bult C.,
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An Wahlestedt C., Mattick J.S., Hume D.A., Kai C., Sasaki D., Tomaru Y.,
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Waki K., Watahiki A., Okamura-Oho Y., Suzuki H., Kawai J.,
Hayashizaki Y.,
Hayashizaki Y., Watahiki A., Okamura-Oho Y., Suzuki H., Kawai J.,
Hayashizaki Y., Watahiki A., Okamura-Oho Y., Suzuki H., Kawai J.,
Hayashizaki Y.,
NUCLEOTILE SPACE NUMBER OF THE NUMBER OF TRAIN=C57BL/6J; TISSUE=Bone marrow;
STRAIN=C57BL/6J; TISSUE=Bone marrow;
MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Kawai J., Shinagawa A., Fukunishi Y., Konno H., Adachi J., Fukuda Arakawa T., Hara A., Fukunishi Y., Konno H., Kondo S., Yamanaka Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka Aizawa K., Izawa M., Nishi K., Bono H., Kasukawa T., Saito
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=C57BL/6J;
PubMed=16141073;
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Science 309:1559-1563(2005).
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d=16141073; DOI=10.1126/science.1112009;

denome Exploration Research Group, and Genome Science Group

me Network Core Team) and the FANTOM Consortium;

sense Transcription in the Mammalian Transcriptome.";

ice 309:1564-1566(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   full-length cDNAs.";
420:563-573(2002).
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Query Match
Best Local
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MGI; MGI:98821; Trf.
G0; G0:0030139; C:endocytic vesicle; IDA.
G0; G0:005768; C:extracellular space; RCA.
G0; G0:000515; C:extracellular space; RCA.
G0; G0:0008199; F:ferric iron binding; RCA.
G0; G0:0006879; P:iron ion homeostas15; RCA.
G0; G0:0006826; P:iron ion transport; RCA.
InterPro; IPR001156; Peptidase_S60.

Pfam; PF00405; Transferrin; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20530913; PubMed=11078861; DOI=10.1101/gr.152600; Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P., Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P., Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M., Suni N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A., Yamamoto R., Matsunoto H., Sakaguchi S., Ikegami T., Kashiwagi K., Fujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M., Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J., Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y., "RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer."; Genome Res. 10:1757-1771(2000).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=C57BL/6J; TISSUE=Bone marrow;
Arakawa T., Carninci P., Fukuda S., Hashizume W., Hayashida K.,
HOri F., Tida J., Imamura K., Imotani K., Itoh M., Kanagawa S.,
Kawai J., Kojima M., Konno H., Murata M., Nakamura M., Ninomiya N.,
Nishiyori H., Nomura K., Ohno M., Sakazume N., Sano H., Sasaki D.,
Shibata K., Shiraki T., Tagami M., Tagami Y., Waki K., Watahiki A.,
Muramateu M., Hayashizaki Y.;
Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=C57BL/6J; TISSUE=Bone marrow;
MEDLINE=20499374; pubMed=11042159; DOI=10.1101/gr.145100;
Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata
Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
"Normalization and subtraction of cap-trapper-selected cDN
prepare full-length cDNA libraries for rapid discovery of
Genome Res. 10:1617-1630(2000).
                                                                                    PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN_1; 1.
PROSITE; PS00206; TRANSFERRIN_2; 2.
PROSITE; PS00207; TRANSFERRIN_3; 2.
SEQUENCE 697 AA; 76698 MW; D3E37B96B856B525 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Functional annotation of Nature 409:685-690(2001).
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                     Similarity
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                     80.0%;
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Score 92; DB
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2; Mismatches
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                                               DB 2;
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RM MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RA MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA HOpkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Altechnko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Carninci P., Prange C.,

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RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Makek J.A., Gunaratne P.H.,

RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      product:transferrin, full insert sequence) (Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:C730037L17 product:transferrin, full insert sequence) (Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:C730048P21 product:transferrin, full insert sequence) (17 days pregnant adult female amnion cDNA, RIKEN full-length enriched library, clone:1920015B11 product:transferrin, full insert sequence) (17 days pregnant adult female amnion cDNA, RIKEN full-length enriched library, clone:1920066F17 product:transferrin, full insert sequence).
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70-FEB-2006, entry version 9.

Transferrin (10 days lactation, adult female mammary gland cDNA, RI.

full-length enriched library, clone:D730004I19 product:transferrin,
full insert sequence) (17 days pregnant adult female amnion cDNA,
full length enriched library, clone:I920024003

RIKEN full-length enriched library, clone:I920024003
                            STRAIN=C57BL/6J;
PubMed=16141072;
                                                                                                                                                                                                                                   STRAIN-C57BL/6J; TISSUE-Amnion, Liver, and Mammary gland; MEDLINE-99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9. Carninci P., Hayashizaki Y.;
                                                                                                                                                                                                      "High-efficiency
                                                                                                                                                                                                                                                                                                                                                                                                                       NIH MGC Project;
Submitted (MAR-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
STRAIN=FVB/N; TISSUE=Mammary tumor.
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                                                                                                               SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences."
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   TISSUE=Amnion, Liver, and Ma
DOI=10.1126/science.1112014;
ukawa T., Katayama S., Gough
                                                                                                                                                                         full-length cDNA
303:19-44(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL/GenBank/DDBJ databases.
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   Gough J., Frith M.C.,
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RA Bajic V.B., Ravasi T., Lenhard B., Wells C., Kodzius R., Shimokawa K.,
RA Bajic V.B., Brenner S.E., Batalov S., Forrest A.R., Zavolan M.,
RA Davis M.J., Wilming L.G., Aidinis V., Allen J.E.,
RA Ambesi-Impiombato A., Apweiler R., Aturaliya R.N., Bailey T.L.,
RA Chiu K.P., Choudhary V., Christoffels A., Clutterbuck D.R.,
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RA Crowe M.L., Dalla E., Dalrymple B.P., de Bono B., Della Gatta G.,
RA Crowe M.L., Dalla E., Dalrymple B.P., de Bono B., Della Gatta G.,
RA Growell-Hemming P., Gingeras T.R., Gojobori T., Green R.E.,
RA Georgi-Hemming P., Gingeras T.R., Gojobori T., Green R.E.,
RA Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa N.,
RA Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa T.,
RA Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa T.,
RA Kitano H., Kollias G., Krishnan S.P., Kruger A., Kummerfeld S.K.,
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RA Matsuda H., Matsuzawa S., Miki H., Mignone F., Miyake S., Morris K.,
RA Misson R., Nishiguchi S., Nishikawa S., Nori F., Ohara O.,
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RA Rost B., Ruan Y., Salzberg S.L., Sandelin A., Schneider C., Pesole G.,
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RA Grimmoda K., Teasdale R.D., Liu E.T., Brusic V., Quackenbush J.,
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RA Wallestedt C., Mattick J.S., Hume D.A., Kai C., Sasaki D., Tomaru Y.,
RA Hayashima T., Kojima M., Kato T., Kawasai H., Kawagashira N.,
RA Hayashirak Y., Nismika N., Nismiya N.,
RA Hayashirak Y., Nismika N., Nismiya N.,
RA Hayashirak M., Nismiya N.,
RA
RA Nazuli Negati Purmed 1, 240001; Pulle 1, 1000 Nacule C. 1200 Na
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=C57BL/6J; TISSUE=Amnion, Liver, and Mammary gland; MBDLINE=22354683; Pubmed=12466851; DOI=10:1038/nature01266; DOI=10:105 H WESTING M WESTING M T NASH; T BOY H W T
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Transcriptome.";
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Gojobori T.,
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RC MEDLINE-21085660; PubMed=11217851; DOI-10.1038/35055500;
RA MEDLINE-21085660; PubMed=11217851; DOI-10.1038/35055500;
RA Marai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Komno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
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RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
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RA Schriml L.M., Staubhi F., Suzuki R., Tomita M., Wagner L., Washio T.,
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RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Scorch K.-F.,
RA Sasaki H., Sato K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Hayashizaki Y.,
RA Hayashizaki Y.,
RA Hayashizaki Y.,
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Best Local S
Matches 18
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Arakawa T., Lina
Hori F., Iida J., Lina
T., Kojima M., K
Nomura j
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Kawai J., Kojim.
Nishiyori H., Nomura
Nishiyori H., Shiraki T
                                                                                                                                                                Nishija
Shibata K., Shi
Smatsu M., J
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MEDLINB-20530913, PubMed=11076861; DOI=10.1101/gr.152600;
Shibata K., Itch M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
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sequencing pipeline with 384 multicapillary sequencer.";
Genome Res. 10:1757-1771(2000).
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Genome R
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Birney E., Hayashizaki Y.;
"Analysis of the mouse transcriptome based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE
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391
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Res. 10:1617-1630(2000).
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420:563-573(2002).
                                                                                    18;
                                                                                                           Similarity
  TTEDCIEKIVNGEADAMTLDGG
                                           TTEDSIAKIMNGEADAMSLDGG
                                                                                                                                                                       Hayashizaki
                                                                                                                                                                ncı P., Fukuda S., Hashizume W., Hayashida K., Imamura K., Imotani K., Itoh M., Kanagawa S., M., Konno H., Murata M., Nakamura M., Ninomiya N., ura K., Ohno M., Sakazume N., Sano H., Sasaki D., ci T., Tagami M., Tagami Y., Waki K., Watahiki A., ishizaki Y.;
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                                                                                  Score 92; DB
Pred. No. 4.5e
2; Mismatches
                                                                                  2
412
                                           22
                                                                                                         DB 2;
4.5e-05;
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                                                                                                                           Length 697;
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RESULT

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Glycoprotein;
Methylation; R
SIGNAL
                                                                                                                                                                                                                                  PRINTS; PR00422; TRANSFER.

SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN 1; 1.

PROSITE; PS00206; TRANSFERRIN 2; 2.

PROSITE; PS00207; TRANSFERRIN 3; 1.

PROSITE; PS00207; TRANSFERRIN 3; 1.
                                            METAL
METAL
METAL
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METAL
BINDING
BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                      --- FUNCTION: Transferrins are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate. It is responsible for the transport of iron from sites of absorption and heme degradation to those of storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.

--- SUBDUIT: Monomer (By similarity)
--- SUBCELLULAR LOCATION: Secreted protein.
--- TISSUE SPECIFICITY: Expressed by the liver and secreted in plasma.
--- MISCELLANEOUS: The peptidase S60 domains do not have protease function as the catalytic residues are missing.
--- SIMILARITY: Belongs to the transferrin family.
--- SIMILARITY: Contains 2 peptidase S60 domains.
                                                                                                                                                                            DOMAIN
                                                                                                                                                                                                                                                                                                                                                                      EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=96132861; PubMed=8557646; DOI=10.1074/jbc.271.2.1166; Retzer M.D., Kabani A., Button L.L., Yu R.H., Schryvers A.B.; "Production and characterization of chieric transferrins for determination of the binding domains for bacterial transferrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    029443;
01-NOV-1997, integrated into U
01-NOV-1997, sequence version
07-MAR-2006, entry version 42.
 BINDING
                                                                                                                                                    METAL
                                                                                                                                                                                                                                                                                                                       PANTHER; PTHR11485; Peptidase
Pfam; PF00405; Transferrin; 2
                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted
Distributed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
Pecora; Bovidae; Bo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 binding globulin)
Name=TF;
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                      BINDING
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                                                                                                                                                                                                                                                                                                                                             InterPro;
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                                                                                                                                                                                                                                                                                                                                                                      U02564; AAA96735.1; P09571; 1H76.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Bovine).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           precursor (Transferrin) (Siderophilin) (Beta-1-metal-
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689
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211
272
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Laurasiatheria;
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heria; Cetartiodactyla;
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TRFL PIG
14632; Q29557;
01-APR-1990, integrated into Uni
15-DEC-1998, sequence version 3
07-FEB-2006, entry version 46.
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           PROTEIN SEQUENCE OF 20-49.

MEDLINE-90105538; PubMed=2605266;
Hutchens T.W., Magnuson J.S., Yip T.-T.;
Harchens T.W., Magnuson J.S., Yip T.-T.;
"Rapid purification of porcine colostral whey lactoferrin by aff chromatography on single-stranded DNA-agarose. Characterization, acid composition and N-terminal amino acid sequence.";
Biochim. Biophys. Acta 999:323-329(1989)
                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE [MRNA].

NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=92379101; PubMede1511016; DOI=10.1016/0167-4781(92)90061-4;

Lvndon J.P., O'Malley B.R., Saucedo O., Lee T., Headon D.R.,
                                                                                                                                                                  NUCLEOTIDE SEQUENCE
MEDLINE=92367939; Po
                                                                                                                                                       Alexander L.J.,
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 FUNCTION:
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J., Levine W.B., Teng
sequencing of the por
23:251-256(1992).
Transferrins
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are iron binding transport proteins which
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 HSSP; Q9TUMO; 1DTZ.

Interpro; IFR001156; Peptidase_S60; 1.

PANTHER; PYHR11485; Peptidase_S60; 1.

Pfam; PF00405; Transferrin; 2.

PFINTS; PR00422; TRANSFERRIN.

SMART; SW00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN 1; 2.

PROSITE; PS00206; TRANSFERRIN 2; 2.

PROSITE; PS00207; TRANSFERRIN 3; 1.

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EMBL; M81327; AAA31059.1;
PIR; A45543; A45543.
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HSSP; Q29477; 1JWI.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60;
PANTHER; PTHR111485; Peptidase_S60; 1.
Pfam; PF00405; Transferrin; 2.
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SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN_1; 2.
PROSITE; PS00206; TRANSFERRIN_2; 2.
PROSITE; PS00207; TRANSFERRIN_3; 1.
SEQUENCE 704 AA; 77612 MW; 10DFEA3C1366D5DA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shinkai H., Morozumi T., Toki D., Muneta Y., Awata T., Uenishi Submitted (JAN-2003) to the EMBL/GenBank/DDBJ databases.
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Eukaryota; Metazoa;
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M -> I (in Ref. 1).
D -> G (in Ref. 1).
Missing (in Ref. 2).
T -> I (in Ref. 1).
E -> S (in Ref. 1).
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By similarity.
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RRT - YTR (in Ref.
M - I (in Ref.
D - G (in Ref.
                                                                                                                     Score 92; DB 2;
Pred. No. 4.5e-05;
3; Mismatches 3
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RESULT 15
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ID Q7YS20_PIG
AC Q7YS20_PIG
AC Q7YS20_PIG
DT 01-CCT-2003
CC Eukaryota; N
OC Mammalia; Ex
OC Mammalia; Ex
OC MAMMILIA
OC Mammalia; Ex
OC NUCLICTIDE S
RP NUCLICOTIDE S
RP NUCLICOTIDE S
RC TISSUE=Mamma
RA PECOTÍNÍ C.,
RL Submitted C.,
CC Copyrighted
CC Copyrighted
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DR HSSP; P24627
DR GO; GO:00066
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Search completed: June 13, Job time : 186.691 secs
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SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN 1; 2.

PROSITE; PS00205; TRANSFERRIN 2; 1.

PROSITE; PS00207; TRANSFERRIN 3; 1.

SEQUENCE 704 AA; 77523 MW; AACBI
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GO; GO:0005576; C:extracellular region; IEA.

GO; GO:0008199; F:ferric iron binding; IEA.

GO; GO:0006879; P:iron ion homeostasis; IEA.

GO; GO:0006826; P:iron ion transport; IEA.
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Pecorini C., Fogher C., Baldi A.;
Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-OCT-2003, integrated into U
01-OCT-2003, sequence version
07-FEB-2006, entry version 11.
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PANTHER; PTHR11485; Peptidase_S60;
Pfam; PF00405; Transferrin; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
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17; Conserv
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Title:

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is derived by analysis of the total score distribution.
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1: /EMC Celerra SIDS3/ptodata/2/iaa/5 COMB.pep:*

2: /EMC Celerra SIDS3/ptodata/2/iaa/6 COMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7 COMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H COMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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US-09-037-188-2

US-09-285-310-2

US-09-753-385-2

US-08-470-058-4

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US-08-1145-681-6
US-08-433-703-6
US-08-456-106-6
US-08-456-108-6
US-09-256-577-6
US-09-633-739-6
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US-09-932-190-2
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  Matches
                    Query Match 95.7%;
Best Local Similarity 95.7%;
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  22;
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ALIGNMENTS

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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/175,158/
FILING DATE: 28-DEC-1993
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/832,029
PRIOR APPLICATION NUMBER: US 07/832,029
PILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
NAME: DECONTA', GAULIO A.
REGISTRATION NUMBER: UVI-005C
TELECHMONICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
FEMORTH: 698 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/08175158A

Patent No. 5986067

Patent No. 5986067

PAPPLICANT: FUNK, Walter D.

APPLICANT: MASON, Anne B.

APPLICANT: MASON, Anne B.

APPLICANT: MODDWORTH, ROBERT C.

APPLICANT: MODDWORTH, RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF

NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESS:
MOLECULE TYPE: protein
                    TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boston
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0; Mismatches

Score 110; DB 1; Pred. No. 1.2e-09;

Length 698; Indels

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REGISTRATION NUMBER: 31,503
REFERENCE/DOCKET NUMBER: UVI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-75941
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 698 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                   Sequence 2, Application US/08470058 Patent No. 5817789 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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APPLICANT:
APPLICANT:
             APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
TITLE OF INVENTION: Chimeric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: WOODWORTH, ROBERT C.
TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF.
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/439,740 FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE:
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                                                                                                                                                                                                                                  392 TTEDCIAKIMNGEADAMSLDGGF 414
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5. 6825037
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60 State Street, suite 510
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                                                                                                                                                                                                                                                                                                        Conservative
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95.7%;
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                                                                                                                                                                                                                                                                                                      Score 110; DB Pred. No. 1.2e 0; Mismatches
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      COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
34,819
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Ph.D., J.D., Janis K.
REGISTRATION NUMBER: 34,819
ATTORNEY/AGENT NUMBER: 34,819
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APPLICANT: Heartl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION UNMBER: 32, 227
REFERENCE/DOCKET NUMBER: TKT9
TELECOMMUNICATION INFORMATION:
TELECHIONE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,058
FILING DATE: 06-JUN-1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                     STREET: 225 F
CITY: Boston
STATE: MA
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CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 02110-2804
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Fish & Richardson P.C.
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o. 6027921
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22; Conserva
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ilarity 95.7%;
Conservative (
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07236/009002
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pred. No. 2e-09;
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TELECOMMUNICATION INFORMATION:

TELEPHONE: 617/542-5070 TELEFAX: 617/542-8906 TELEX: 200154

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US-09-285-310-2
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                                                                Matches
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COMPUTER READABLE FORM:
COMPUTER: Diskette
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
OPERATING SYSTEM: Windc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1074 amino aci
                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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NAME: Fraser, Ph.D., J.D., J
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 072
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APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/285,310
                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
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                                                                                                                                                                              TYPE:
                                                                                                                                                                                                                                             TELEPHONE: 617/542-50
TELEFAX: 617/542-8906
TELEX: 200154
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                                                                               Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        768 TTEDCIAKIMNGEADAMSLDGGF 790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
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                                                                               Similarity
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TTEDCIAKIMNGEADAMSLDGGF 790
                   TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MA
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225 Franklin Street
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SYSTEM: Windows95
                                                                                                                                               protein
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95.7%;
                                                                             95.7%;
95.7%;
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                                                              0; Mismatches
                                                                             Score 110; DB 2
Pred. No. 2e-09;
                                                                                                                                                                                                                                                                                                                  07236/009002
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Pred. No. 2e-09;
                                                                                                                                                                                                                                                                                                                                               Janis K.
                                                                                        DB 2; Length 1074;
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US-09-753-385-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/08470058 Patent No. 5817789
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/470,058
FILING DATE: 06-JUN-1995
CLASSIFICATION: 530
CLASSIFICATION: 530
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PRIOR FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1998-03-09
PRIOR APPLICATION NUMBER: US 09/285,310
PRIOR FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Conclio, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE IN TRANSPORTITE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS
FILE REFERENCE: 10278-025004
CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
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TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                               CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REGISTRATION NUMBER: TK7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey Fr
TITLE OF INVENTION: Chimeric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
                                            REFERENCE/DOCKET NUMBER: TK
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Two Mil:
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02173
                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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Two Militia Drive
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ilarity 95.7%;
Conservative
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                                                                                             TKT93-01
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Pred. No. 2e-09;
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                                                                                                                                                                                                                                                                                   Version #1.30
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RESULT 9
US-09-285-310-4
; Sequence 4, Application US/09285310
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                                                                                                                                                                                                         ; MOLECULE TYPE: protein US-09-037-188-4
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Best Local (
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Best Local
                                                                                                                                                                                                                                                                                   TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                  NAME: Fraser, Ph.D., J.D., Janis K. REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07236/009002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 06-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 02-MAR-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: CHI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible OPERATING SYSTEM: Window
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 225 (CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: un
TOPOLOGY: unknow
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                                                                                                                                                                                                                                         TYPE: amino acid TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                   TELEFAX: 617/542-8906
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                                                                                  1104 TTEDCIAKIMNGEADAMSLDGGF 1126
                                                                                                                                             22;
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Similarity 95.7%;
22; Conservative
                                                                                                                                                             Similarity
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                                                                                                                                             Conservative
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                                                                                                                                                          95.7%;
95.7%;
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                                                                                                                                                           Score 110; DB 2;
Pred. No. 2.7e-09;
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Pred. No. 2.7e-09;
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APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
APPLICANT: Concino, Michael F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE IN TRANSPORT TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS FILE REFERENCE: 10278-025004
CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: US 09/037,188
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/09753385
Patent No. 6858578
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APPLICANT: Heartl
APPLICANT: Lemont
                    NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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INFORMATION FOR SEQ ID NO:
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FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: Past-SEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unknown MOLECULE TYPE: protein
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LENGTH: 1410 amino aci
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REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
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APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHYMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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CITY: Boston
STATE: MA
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NO 4
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                       for Windows Version 4.0
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Pred. No. 2.7e-09;
0; Mismatches 1
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RESULT 11
5262177-4
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Best Local Similarity
Gardhes 22; Conservi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-753-385-4
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US-09-724-864-54
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                                                                                                                                        ; ORGANISM: Mouse
US-09-724-864-54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Patent No. 5262177
; APPLICANT: BROWN, J OSEPH P.;ESTIN, CHARLES D.;PLOWMAN, GREGORY
; DI;HELLSTROM, KARL E.;ROSE, TIMOTHY M.;HELLSTROM, INGEGERD;
; DURCHIO, ANTHONY F.;HU, SHIU-LOK;PENNATHUR, SRIDHAR
; TITLE OF INVENTION: RECOMBINANT VIRUSES ENCODING THE HHUMAN
;MELANOMA-ASSOCIATED ANTIGEN
                                                                                                                                                                        SOFTWARE: Fa
SEQ ID NO 54
LENGTH: 697
TYPE: PRT
                                                                                                                                                                                                                                                                                Patent No. 6380362

GENERAL INFORMATION:

APPLICANT: Watson, James D

APPLICANT: Murison, James G.

TITLE OF INVENTION: Polynucleotides, polypeptides expressed

TITLE OF INVENTION: by the polynucleotides and methods for their use.

FILE REFERENCE: 11000.1050U1

CURRENT APPLICATION NUMBER: US/09/724,864

CURRENT FILING DATE: 2000-11-28

CURRENT FILING DATE: 2000-11-28
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Matches
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                                                                                                    Query Match
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/7230
FILING DATE: 27-JAN-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 827,313
FILING DATE: 07-FEB-1986
                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: U.S. NO. 6380362 60/171,678
PRIOR FILING DATE: 1999-12-23
NUMBER OF SEQ ID NOS: 72
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nes 21; Conserv
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                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          392
 391
                                                                                    Similarity
TTEDCIEKIVNGEADAMTLDGG 412
                                  TTEDSIAKIMNGEADAMSLDGG 22
                                                                                                                                                                                                                           FastSEQ
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ilarity 91.3%;
Conservative
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95.7%;
                                                                                  80.0%;
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; Pred. No. 6.3e
0; Mismatches
                                                                  Score 92; DB 2; Le
Pred. No. 1.1e-06;
2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 110; DB 2;
Pred. No. 2.7e-09;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 7;
                                                                                                  Length 697;
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RESULT 14
US-08-453-703-6
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US-08-145-681-6
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                                                                                                                                                                                                              Sequence 6, Application US/08453703 Patent No. 5766939
                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/145,681
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MCGTeggor, Martin L.
REGISTRATION NUMBER: 29,329
REGISTRATION NUMBER: 29,329
REFERENCE/DOCKET NUMBER: 19928-012:
                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 713/229/1522
INFORMATION FOR SEQ ID NO:
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APPLICANT: May, Greg
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 77002
ZIP: 77002
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                              APPLICANT: May, Gregory S.

TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using CDI
TITLE OF INVENTION: Various Organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
ORGANISM: Sus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein HYPOTHETICAL: YES ANTI-SENSE: NO
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 703 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
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es 17; Conserv
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910 Louisiana St
                                                                                                                                      Conneely, Orla M. Headon, Denis R. O'Malley, Bert W.
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                                                                                        and Lactoferrin Polypeptides Using cDNA Sequences in
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Pred. No. 1.1e-0
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CITY: New York STATE: New York COUNTRY: USA

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US-08-456-106-6
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ORGANISM: Sub scrofa
US-08-453-703-6
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TELEPHONE: 415-854-3660
TELEPAX: 415-854-3694
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 703 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/08456106 Patent No. 5849881 GENERAL INFORMATION:
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Best Local (
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APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using cDNA Sequences in
TITLE OF INVENTION: Various Organisms
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & FAmerical Penn
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FILING DATE: Concurrently herewith CLASSIFICATION: 435
PRIOR APPLICATION DATA: 08/145,681
APPLICATION NUMBER: 08/145,681
FILING DATE: October 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCY/DOCUMENTER: 25,227
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ANTI-SENSE: NO
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/456,106
FILING DATE: Concurrently herewith
CLASSIFICATION: 435
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                  STATE: N
                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                      ZIP: 10036
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390 TTEDCIVQVLKGEADAMSLDGGF 412
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 Mismatches

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Pred. No. 1.1e-06;
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; ORIGINAL SOURCE:
; ORGANISM: Sus
US-08-456-106-6
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                                                          Query Match 80.0
Best Local Similarity 73.9
Matches 17; Conservative
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                   MOLECULE TYPE: protein
HYPOTHETICAL: YES
ANTI-SENSE: NO
                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 703 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                  TELEPHONE: 415-854-3660
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,68:
FILING DATE: 0Ctober 28, 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8206-025
390 TTEDCIVQVLKGEADAMSLDGGF 412
                     TTEDSIAKIMNGEADAMSLDGGF 23
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                                                             3; Mismatches
                                                                          Score 92; DB 1;
Pred. No. 1.1e-06
                                                                                           Length 703;
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Search completed: June 13, 2006, 04:25:49 Job time : 40.5882 secs

Gaps

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Result
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Maximum DB seq
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Perfect score:
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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length: 2000000000
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:*
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115
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.
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US-09-891-126-5
US-10-266-745-5
US-10-378-094-3
US-10-314-94-3
US-10-429-482-4
US-10-429-515-4
US-10-429-535-4
US-10-429-635-4
US-10-429-655-4
US-10-429-660-4
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US-10-429-660-4
US-10-429-661-4
US-10-429-662-4
US-10-429-662-4
US-10-429-655-4
US-10-429-655-4
US-10-318-094-2
US-10-318-094-2
US-10-318-094-2
US-10-318-201-10
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ALIGNMENTS	US-10-169-297-9	US-10-987-587-2	US-10-170-221-2	US-10-205-331-55	US-10-316-253-275	US-10-316-253-273	US-10-023-096-2	US-10-639-835-4	US-10-077-381-4	US-10-076-816-4	US-10-169-297-49	US-10-513-523-4	US-10-620-256-6	US-10-473-127-804	US-10-473-127-805	US-09-753-385-4	US-09-753-385-2	US-11-038-901-2	
	Sequence 9, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 55, Appl	Sequence 275, App	Sequence 273, App	Sequence 2, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 49, Appl	Sequence 4, Appli	Sequence 6, Appli	Sequence 804, App	Sequence 805, App	Sequence 4, Appli	Sequence 2, Appli	Sequence 2, Appli	

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Sequence 2, Application US/10612162

Publication No. US20040014145A1

GENERAL INFORMATION:
APPLICANT: Dade Behring Marburg GmbH
ITITLE OF INVENTION: antibodies, their preparation and use
FILE REFERENCE: 2002/B001

CURRENT APPLICATION NUMBER: US/10/612,162

CURRENT FILING DATE: 2003-07-03

PRIOR APPLICATION NUMBER: 10230550.1

PRIOR APPLICATION NUMBER: 10230550.1

PRIOR APPLICATION NUMBER: 0303-05-19

NUMBER OF SEQ ID NOS: 4

SOFTWARE: Patentin Ver. 2.1
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US-09-891-126-5
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US-10-612-162-2
Sequence 5, Application US/09891126
Patent No. US2002072596A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Transferrin Polynucleotides, Polypeptides, and Antibodies
FILE REFERENCE: PT035P1
CURRENT APPLICATION NUMBER: US/09/891,126
CURRENT FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR FILING DATE: 1999-12-23
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LENGTH: 23
TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver.
SEQ ID NO 5

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Sequence 3, Application US/10378094
Publication No. US20030221201A1
GENERAL INFORMATION:
APPLICANT: PRIOR, Christopher P.
APPLICANT: LAI, Char-Huei
APPLICANT: SADEGHI, Homayoun
APPLICANT: NUMBER: US/10/378,094
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT FILING DATE: 2003-03-04
PRIOR APPLICATION NUMBER: US/031,494
PRIOR APPLICATION NUMBER: US/031,494
PRIOR APPLICATION NUMBER: US/0334,059
PRIOR APPLICATION NUMBER: US/0334,059
PRIOR APPLICATION NUMBER: US/0334,059
PRIOR FILING DATE: 2001-11-30
PRIOR FILING DATE: 2001-10-30
PRIOR FILING DATE: 2001-10-30
PRIOR FILING DATE: 2001-10-30
PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 66
SOFTMARE: Patentin version 3.2
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US-10-266-745-5
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CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US/09/891,126
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR FILING DATE: 2000-12-21
PRIOR PPLICATION NUMBER: 60/171,595
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR FILING DATE: 1999-12-23
NUMBER OF SEQ ID NOS: 7
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Publication No. US20030149256A1
GENERAL INFORMATION:
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                                                                                                                 SEQ ID NO 3
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TITLE OF INVENTION: Transferrin Polynucleotides, Polypeptides, and Antibodies
FILE REFERENCE: PT035P1
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ORGANISM: Homo sapiens
                                        ORGANISM: Homo sapiens
                                                                         TYPE: PRT
NAME/KEY: MISC_FEATURE
                                                                                       ENGTH:
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95.7%;
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Pred. No. 8.4e-09;
0; Mismatches 1
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Pred. No. 8.4e-09
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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US-10-231-494-3
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PRIOR FILING DATE: 2002-08-30
PRIOR PELICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
                                                                                                                                                                                      Sequence 3, Application US/10231494
Publication No. US20040023334A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
CURRENT APPLICATION NUMBER: US/10/231,494
CURRENT APPLICATION NUMBER: US/0/231,745
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2001-08-30
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
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Best Local Similarity
Matches 22; Conserv
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TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
FILE REFERENCE: 54710-5004-US
CURRENT APPLICATION NUMBER: US/10/384,060
CURRENT FILING DATE: 2003-03-10
CURRENT FILING DATE: 2003-03-10
                                                                                                                              NUMBER OF SEQ ID NOS: 30
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ORGANISM: Homo sapiens
FEATURE:
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OTHER INFORMATION: Mature Transferrin Protein
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nes 22; Conser
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Pred. No. 2e-08;
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Pred. No. 2e-08;
0; Mismatches
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US-10-429-497-4
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GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For The Treatment Of Tumors
                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/10429497
Publication No. US20040219098A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For The Treatment
FILE REFERENCE: 2537.000006
CURRENT APPLICATION NUMBER: US/10/429,497
CURRENT FILING DATE: 2003-05-02
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Publication No. US20040219097A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The Diagnosis, Imaging and Treatment Of FILE REFERENCE: 2537.000001
CURRENT APPLICATION NUMBER: US/10/429,482
CURRENT FILING DATE: 2003-05-02
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Best Local Similarity 95.7%;
Matches 22; Conservative
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SEQ ID NO 4
LENGTH: 679
TYPE: PRT
                                                Sequence 4, Application US/10429515 Publication No. US20040219099A1
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SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 6
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ORGANISM: Homo sapiens
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mes 22; Conservative
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95.7%;
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Pred. No. 2e-08;
0; Mismatches
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Pred. No. 2e-08;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Pred. No. 2e-08;
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-429-635-4
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-429-598-4
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US-10-429-635-4
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                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1 SEQ ID NO 4
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                   TTEDCIAKIMNGEADAMSLDGGF 395
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CURRENT APPLICATION NUMBER: US/10/429,515
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOPTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
US-10-429-515-4
                                                                                                                                                                                Sequence 4, Application US/10429635
Publication No. US20040219101A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For Treatment Of Tumors
FILE REFERENCE: 2537,000007
CURRENT APPLICATION NUMBER: US/10/429,635
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The Treatment of Tumors
FILE REFERENCE: 2537.000003
CURRENT APPLICATION NUMBER: US/10/429,598
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEG ID NOS: 6
SOFTWARE: Patentin version 3.1
SEG ID NO 4
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Pred. No. 2e-08;
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APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Compositions For Drug Delivery
FILE REFERENCE: 2537.000009
CURRENT APPLICATION NUMBER: US/10/429,653
CURRENT FILING DATE: 2003-05-02
UNMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
LENGTH: 679
TYDE: PRT
ORGANISM: Homo sapiens
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Publication No. US20040219103A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods Useful For The Diagnosis, Imaging and Treatment Of Tumore
CURENT APPLICATION UMMBER: US/10/429,659
CURRENT APPLICATION UMMBER: US/10/429,659
CURRENT FILING DATE: 2003-04-30
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
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US-10-429-653-4
US-10-429-661-4
                                APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For Treatment Of Tumors
FILE REFERENCE: 2537.000008
CURRENT APPLICATION NUMBER: US/10/429,661
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
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Best Local Similarity 95.7%;
Matches 22; Conservative
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Best Local Similarity
                ORGANISM: Homo sapiens
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GENERAL INFORMATION:

APPLICANT: Sandhu, Jasbir

ITILE OF INVENTION: Methods For Nucleic Acid Delivery

FILE REFERENCE: 2537.000012

CURRENT APPLICATION NUMBER: U5/10/429,660

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 12

SOFTWARE: PatentIn version 3.1

SEQ ID NO 4

LENGTH: 679

Type: PAT

ORGANISM: Homo sapiens

US-10-429-660-4
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Search completed: June 13, Job time : 132.25 secs
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Publication No. US20040220084A1
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Local Similarity 95.7%;
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Listing first 45 summaries
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1: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US09_NEW_PUB.pep:*

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3: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

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8: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
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US-11-258-767-27

US-11-258-767-20

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83 72.2 695 7 US-11-258-767-26 82 71.3 711 7 US-11-258-767-34 77 67.0 681 7 US-11-258-767-13 77 67.0 686 6 US-10-515-429-43 77 67.0 708 7 US-11-258-767-22 77 67.0 708 7 US-11-258-767-22 77 67.0 708 7 US-11-258-767-29 78 70 708 7 US-11-258-767-39 79 67.0 708 7 US-11-258-767-39 70 67.0 708 7 US-11-258-767-39 71 67.0 708 7 US-11-258-767-39 72 67.0 708 7 US-11-258-767-39 73 39.1 325 6 US-10-520-999-7 745 39.1 325 6 US-10-520-999-7 75 39.1 325 6 US-10-520-999-7 76 39.1 336 US-10-520-999-7 77 67.0 708 7 US-11-258-767-39 78 39.1 316 US-10-520-999-7 79 67.0 708 7 US-11-258-767-39 79 67.0 708 7 US-11-258-767-39 70 708 7 US-11-258-767-39 70 67.0 708 7 US-11-258-767-39 71 67.0 708 7 US-11-258-767-39 72 67.0 708 7 US-11-258-767-39 73 39.1 325 6 US-10-520-999-1 74 38.3 1100 6 US-10-520-999-1 75 13 13 13 13 13 13 13 13 13 14 15 15 15 15 15 15 15 15 15 15 15 15 15	72.2 695 7 US 71.3 711 7 US 67.0 686 6 US 67.0 708 7 US 67.0 333 6 US 39.1 325 6 US 39.1 371 6 US 39.1 530 6 US 39.1 530 6 US 39.1 530 6 US	45	44	4 W	42	41	40	39	38	37	36	35	ω 4	ω ω	32	31	30	29	28	27	
2 695 7 US 3 711 7 US 6 681 7 US 6 686 6 US 6 6 6 US 6 708 7 US 7 7 7 US 7 US	2 695 7 US-11-258-767-26 2 695 7 US-11-258-767-34 3 711 7 US-11-258-767-34 5 equence 6 681 7 US-11-258-767-13 6 equence 7 686 6 US-10-515-429-43 6 equence 7 687 7 US-11-258-767-16 7 08 7 US-11-258-767-26 8 equence 7 7 US-11-258-767-27 8 equence 7 7 US-11-258-767-28 8 equence 7 7 US-11-258-767-29 9 equence 7 7 US-11-258-767-39 9 equence 7 7 US-11-258-767-39 9 equence 7 7 US-11-258-767-39 9 equence 1 325 6 US-10-259-99-1 1 331 6 US-10-520-999-1 1 342 6 US-10-520-999-1 1 353 6 US-10-520-999-8 1 151 6 US-10-5	44	44	45	45	45	45	45	45	77	77	77	77	77	77	77	77	77	82	83	•
6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	7 US-11-258-767-26 7 US-11-258-767-34 Sequence 7 US-11-258-767-13 Sequence 8 US-10-258-767-13 Sequence 9 US-11-258-767-13 Sequence 10 US-11-258-767-16 Sequence 10 US-11-258-767-22 Sequence 10 US-11-258-767-23 Sequence 10 US-11-258-767-29 Sequence 10 US-11-258-767-35 Sequence 10 US-11-258-767-35 Sequence 10 US-11-258-767-35 Sequence 10 US-11-258-767-39 Sequence 10 US-10-520-999-10 Sequence 10 US-10-520-999-10 Sequence 10 US-10-520-999-10 Sequence 10 US-10-520-999-10 Sequence 10 US-10-515-429-63 Sequence 10 US-10-515-439-63 Sequence 10 US-10-93-349-1941 Sequence 10 US-10-93-349-1941	38.3	38.3	39.1	39.1	39.1	φ	39.1					67.0	67.0							
7 US-11-258-767-34 7 US-11-258-767-34 7 US-11-258-767-34 7 US-11-258-767-16 US-10-515-429-43 7 US-11-258-767-22 7 US-11-258-767-22 7 US-11-258-767-29 7 US-11-258-767-39 7 US-11-258-767-39 8 US-10-520-999-10 9 US-10-520-999-10 9 US-10-520-999-9 9 US-10-520-999-9 9 US-10-520-999-12	-11-258-767-26 -11-258-767-34 -11-258-767-33 -11-258-767-13 -11-258-767-13 -11-258-767-13 -11-258-767-13 -11-258-767-22 -11-258-767-22 -11-258-767-29 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-258-767-39 -11-25	1151	1100	530	451	371	342	333	325	708	708	708	708	708	708	708	686	681	711	695	
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ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
ITILE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2003-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 679
                                                                                                                                                                                                                                                                 ; FEATURE:
; OTHER INFORMATION: Mature transferrin protein
US-10-515-429-3
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US-10-515-429-3
US-10-515-429-3
Sequence 3, Application US/10515429
Publication No. US20060105387A1
RESULT 2
US-10-515-429-2
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                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo sapiens
                                                                                     373
                                                                                                          1 TTEDSIAKIMNGEADAMSLDGGF 23
                                                                                                                                                                         l Similarity
22; Conserv
                                                                                   TTEDCIAKIMNGEADAMSLDGGF 395
                                                                                                                                                                           Conservative
                                                                                                                                                                                            95.7%;
95.7%;
                                                                                                                                                                           0;
                                                                                                                                                                                                Score 110; DB 6;
Pred. No. 1.4e-09;
                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                     Length 679;
                                                                                                                                                                              Indels
                                                                                                                                                                              0
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Sequence 2, Application US/10515429
Publication No. US20060105387A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries

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RESULT 4
US-10-515-429-40
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                                                                                                                                                                                                                                                                      LENGTH: 676
; TYPE: PRT
; ORGANISM: Oryctolagus cuniculus
US-10-515-429-37
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CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
SOFTWARE: PACENTIN Ver. 2.1
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Sequence 40, Application US/10515429
Publication No. US20066105387A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 37
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LENGTH: 698
                                                                                                                                                                                                               Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                        373 TPEDCIAKIMNGEADAMSLDGGY 395
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22; Conservat
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                                                                                                                                                                      1 TTEDSIAKIMNGEADAMSLDGGF 23
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                                                                                                                                                                                                                             87.8%;
                                                                                                                                                                                                                 1;
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                                                                                                                                                                                                               Score 101; DB 6; Length 676; Pred. No. 3.4e-08; 1; Mismatches 2; Indels
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Pred. No. 1.4e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
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RESULT 6 US-10-515-429-41

Sequence 41, Application US/10515429 Publication No. US20060105387A1 GENERAL INFORMATION:

APPLICANT: Prior, Christopher P. APPLICANT: Turner, Andrew J.

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APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-03-09
PRIOR FILING DATE: 2003-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TITLE OF INVENTION: Transferrin Fusion Protein Libraries
; FILS REFERENCE: 054710-5007-W0
; CURRENT APPLICATION NUMBER: US/10/515,429
; CURRENT FILING DATE: 2004-11-23
; PRIOR APPLICATION NUMBER: PCT/US03/26779
; PRIOR FILING DATE: 2003-08-26
; PRIOR FILING DATE: 2003-08-26
; PRIOR APPLICATION NUMBER: US 60/406,977
; PRIOR FILING DATE: 2002-08-30
; PRIOR APPLICATION NUMBER: US 10/384,060
; PRIOR APPLICATION NUMBER: US 10/384,060
; PRIOR APPLICATION NUMBER: US 10/384,060
; PRIOR APPLICATION NUMBER: US 0/485,404
; PRIOR FILING DATE: 2003-07-09
; NUMBER OF SEQ ID NOS: 75
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                                                                                                                                                                                                 ; TYPE: PRT ; ORGANISM: Mus musculus US-10-515-429-39
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US-10-515-429-39
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; TYPB: PRT
; ORGANISM: Equus caballus
US-10-515-429-40
                                                                                            Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 75
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 39
LENGTH: 677
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 83.5%;
Best Local Similarity 82.6%;
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          376 STEDCIAKIVKGEADAMSLDGGF 398
372 TTEDCIEKIVNGEADAMTLDGG 393
                            1 TTEDSIAKIMNGEADAMSLDGG 22
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No. US20060105387A1
                                                                                                  Conservative
                                                                                                                         80.0%;
                                                                                                  2; Mismatches
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Pred. No. :
                                                                                                                         Score 92; DB 6;
Pred. No. 8.4e-07;
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                                                                                                                                                Length 677;
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LENGTH: 686
; TYPE: PRT
; ORGANISM: PIG
US-11-258-767-36
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US-10-515-429-42
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US-10-515-429-41
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PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
                                                      Sequence 42, Application US/10515429
Publication No. US20060105387A1
GENERAL INFORMATION:
APPLICANT: Frior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.3
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 36, Application US/11258767 Publication No. US20060094082A1
                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
CURRENT APPLICATION NUMBER: US/10/515,429
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                                   PPLICANT: Turner, Andrew J.
PPLICANT: Sadeghi, Homayoun
ILLE OF INVENTION: Transferrin Fusion Protein Libraries
                                                                                                                                                                                                                                                                                                                                                              Local
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Pred. No. 8.5e-07,
3; Mismatches
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Pred. No. 8.5e-07;
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; SOFTWARE: PatentIn v

; SEQ ID NO 37

; LENGTH: 703

; TYPE: PRT

; ORGANISM: PIG

US-11-258-767-37
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US-11-258-767-37
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                                                                                                                      Sequence 38, Application US/10515429 Publication No. US20060105387A1 GENERAL INFORMATION:
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PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 42
SEQ ID NO 42
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Best Local Similarity
Matches 17; Conserv
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APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
CURRENT APPLICATION NUMBER: US/10/515,429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HOO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR TILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Varadhachary, Atul APPLICANT: Glynn, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (308)..(308)
OTHER INFORMATION: Xaa can be any naturally occurring amino
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ORGANISM: Sus scrofa
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Pred. No. 8.7e-07;
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Pred. No. 8.8e-07;
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                                                                                                                                                                                                                                                                                                                                                                                           Length 703;
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Sequence 25, Application US/11258767

Publication No. US20060094082A1

GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2006-10-26
RUYMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.3
SEQ ID NO 25
LENGTH: 698
TYPE: PRT
ORGANISM: RAT
US-11-258-767-25
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US-11-258-767-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-07-09
Sequence 31, Application US/11258767

Publication No. US20060094082A1

GENERAL INFORMATION:

APPLICANT: Varadhachary, Atul

APPLICANT: Petrak, Karel

APPLICANT: Engelmayer, Jose

TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF

FILE REFERENCE: HO-P02915US1

CURRENT FILING DATE: 2005-10-26

PRIOR APPLICATION NUMBER: US 60/622,176

PRIOR FILING DATE: 2004-10-26
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Matches
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SEQ ID NO 38
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PRIOR APPLICATION NUMBER: PCT/US03/26779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 78.3%;
Local Similarity 81.8%;
les 18; Conservative
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81.8%;
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Pred. No. 1.8e-06;
2; Mismatches 2;
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Pred. No. 1.7e-06;
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; NUMBER OF SEQ ID NOS: 67
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 31
; SEQ ID NO 31
; LENGTH: 709
; TYPE: PRT
; ORGANISM: HUMAN
US-11-258-767-31
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US-11-258-767-12
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Best Local Similarity 73.>
""" Conservative
                                                 SOFTWARE: PatentIn version 3.3
SEQ ID NO 12
LENGTH: 711
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 33
LENGTH: 710
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                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                     Sequence 12, Application US/11258767 Publication No. US20060094082A1
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APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
APPLICANT: Engelmayer, Jose
APPLICANT: Engelmayer, Jose
FILE REFERNCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
REGRESSING DATE: 2004-10-26
                                                                                                       CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
                                                                                                                                                                                                      APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
                 TYPE: PRT
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
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No. US20060094082A1
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73.9%;
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3; Mismatches

 Mismatches

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Pred. No. 1.8e-06;
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Query Match

78.3%; Score 90;

DB 7;

Length 711;

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RESULT 15
US-11-258-767-15
Sequence 15, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
SEQ ID NO 15
LENGTH: 171
CURRENT FILING DATE: 2004-10-26
NUMBER: US/11/258-767-15
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Search completed: June 13, 2006, 04:56:53 Job time : 7.77941 secs
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SUMMARIES

Result

Query

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ALIGNMENTS

ADG46076 standard; peptide;

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RESULT 1
ADG46076
ID ADG4
XX ADG4
XX ADG4
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XX ATM
DT 25-M
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DT 19-M
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XX DF 13 05-JUL-2002; 2002DE-01030550. 19-MAY-2003; 2003EP-00011334. EP1378521-A1. antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. Human CDT peptide fragment #2. 25-MAR-2004 ADG46076; 07-JAN-2004. Homo sapiens (DADE-) DADE BEHRING MARBURG GMBH. (first entry)

Claim 4; SEQ ID NO 2; 21pp; German.

New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization.

WPI; 2004-073743/08.

Althaus H;

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSWGKEDLIWELL, TTEDSIAKIMNGRADAMSLDGGF, SKLSMGSGLNLSEEN and YEKYLGEEYVKAN (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for

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RESULT 2
AEE39174
ID AEE3
The invention relates to identifying glycopolypeptides in a serum or CC plasma sample comprising immobilizing derivatized sample glycopeptides to a solid support, releasing the sample glycopeptide CC glycopolypeptides to a solid support, adding to the released sample CC glycopeptide fragments standard peptides, and identifying released sample CC glycopeptide fragments that correspond to standard peptides added by mass gentroscopy. Also included are a method for identifying one or more CC diagnostic markers for a disease, a composition comprising peptides accountaining the glycopylation sites (AEE35897-AEE339378, where the peptides CC cancaining the glycopylation sites (AEE35897-AEE339378). The methods are useful for the used for blood serum profiling for the detection of prognostic and CC diagnostic protein markers. It can also be used to identify and/or validate drug targets and to evaluate drug efficacy, drug dosing, and/or
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Matches 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 3278; 193pp; English.
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Query Match Best Local Similarity

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                                                 The invention relates to a novel method for the quantitative analysis of a protein. The method involves hydrolyzing a first test sample in water containing 90% or more of 18-0 (an isotopic labeling element), hydrolyzing a second sample in water containing 90% or more of 16-0, mixing the hydrolyzed substance of the first and second samples, and carrying out quantitative analysis of the liquid mixture by liquid chromatography/electrospray ionization-mass spectrometry (LC/ESI-MS). The method enables the quantification of proteins in large-scale proteomics, and trace amount of proteins present in an organism can be quantified accurately and efficiently. This sequence represents a peptide fragment used in the protein quantitative analysis method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           drug toxicity. The methods can also be used for the detection of changes in the state of glycosylation of proteins based on the concurrent application of protein abundance measurement of protein glycosylation on the same sample. The method allows fast throughput and simplicity. It can be readily adapted for high throughput analysis of samples, which can be particularly advantageous for the analysis of clinical specimens. The method can also be automated to facilitate the processing of multiple samples. The present sequence is a human glycopeptide that does not contain an N-linked glycosylation site, suitable for use as a reference peptide in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Quantitative analysis of protein, by hydrolyzing test sample and control in water containing labeled oxygen, mixing hydrolyzed substance and subjecting liquid mixture to liquid chromatography/electrospray
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page; 13pp; Japanese.
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SUMITOMO CHEM CO
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Å
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9.3e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of identifying a protein comprising receiving mass spectrometry data that includes putative proteins, peptides and peptide confidence value, identifying and grouping proteins that share peptide information and determining for each group if it is likely that only one protein is present or if peptide evidence exists to support that other proteins in the group are present. The invention is used for identifying proteins in the field of drug discovery, in protein and gene expression analysis, as biomarkers and in other areas of systems biology. This sequence represents a peptide used in the scope of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identification of protein for e.g. drug discovery involves grouping proteins based on mass spectrometry data and determining whether only one form of protein is present or there is evidence to support the presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seymour S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-2003; 2003US-0531405P.
05-AUG-2004; 2004US-0599321P.
                             01-DEC-2005
                                                                                                                                           AEC83228 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mass spectroscopy; gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic peptide #80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-SEP-2005
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                                                                                     AEC83228;
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                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 54.1
                                                                                                                                                                                                                                                                                                                                           11 NGEADAMSLDG 21
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                          (first entry)
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                                                                                                                                        peptide; 23 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                           Score 32; DB 9;
Pred. No. 9.4e+02;
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RESULT 6 ADV12367

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ADV12367 standard; peptide; 23 AA.

ADV12367;

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NGEADAMSLDG 21 NGETTTQGLDG 19

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       composition comprising the sperm flagellar energy carrier protein, and a carrier; an isolated nucleic acid encoding the sperm flagellar energy carrier protein; a vector comprising the nucleic acid; a host cell comprising the vector; a composition comprising an isolated nucleic acid complementary to a nucleic acid encoding a sperm flagellar energy carrier protein, and a carrier; an antibody that binds to a sperm flagellar energy carrier protein; a pharmaceutical composition comprising the antibody and a carrier; an antigenic composition comprising a protein having SEQ ID NO: 2; and a carrier; a method of diagnosing a sperm flagellar energy carrier protein-associated disease or disorder related to aberrant sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein-associated disease or disorder; a method of regulating sperm flagellar energy carrier protein expression, function, or levels in a
                                                                                                                                                                                                                                                                                                                                flagellar energy carrier protein expression, function, or levels in a subject. The sperm flagellar energy carrier protein is useful for treating sperm flagellar energy carrier protein-associated disease or disorder, e.g. infertility. The protein is also useful as contraceptive agent. The present sequence is a peptide derived from human Aldolase A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The new invention relates to a new mammalian sperm specific fibrous sheath protein, that is believed to function as an adenine nucleotide translocase, and has been designated sperm flagellar energy carrier (SPEC) protein comprising SEQ ID No: 2 or 4. The protein is located in the principle piece of the sperm tail, but not in the midpiece. SFEC is believed to be essential for sperm motility, and thus antagonists of SFEC may have use as contraceptive agents. Also claimed are a pharmaceutical
                                                                                                                                                                                            Sequence 23 AA;
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30-SEP-2004; 2004US-0614817P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 8; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New mammalian sperm flagellar energy carrier (SPEC) protein, its homolog, derivative or fragment, useful as contraceptive, and for treating sperm flagellar energy carrier protein-associated disease or disorder, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2005-649557/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kim Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYVI-) UNIV VIRGINIA PATENT FOUND
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                                                                                                                                                                                                                                                                                            fructose-bisphosphate
     Conservative
                                             27.8%;
     <u>,</u>
                                                  Score 32; DB 9;
Pred. No. 9.4e+02;
Mismatches
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10-MAR-2005 (first entry)

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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                               acid sequence but which is not phosphorylated), a system (comprising a computer memory comprising data files storing information relating to the identifying characteristics of positively charged peptides, and a data analysis module capable of executing instructions for organizing and/or searching the data files), a computer program product (comprising data relating to the identifying characteristics of positively charged peptides and comprising instructions for organizing and/or searching the data), and a method for identifying N-terminal peptides in a sample. The present sequence is a peptide from a human phosphoprotein, containing a phosphorylation site, identified by the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     isolated polypeptide selected from any of the polypeptides listed in the specification and is modified at a modification site, an isolated peptide comprising a mass spectral peak signatures. A method for identifying a treatment that modulates phosphorylation of an amino acid in a target polypeptide, a method for generating a peptide standard, a pair of peptide standards comprising the peptide obtained (where the peptide is phosphorylated and a corresponding peptide comprising an identical amino phosphorylated and a corresponding peptide comprising an identical amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (comprising determining the presence, absence or level of one ore more phosphorylated peptides as identified above in cells having a cell state and determining the degree of correlation between the presence, absence or level of phosphorylated polypeptide with the cell state), an isolated peptide of 5-50 amino acids comprising an amino acid sequence that is a subsequence of any of the protein sequences given in the specification (and which comprise a phosphorylation site within the subsequence), an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to characterizing phosphorylated polypeptides in a sample comprising digesting the polypeptides with a protease thus generating test peptides, and collecting a fraction of test peptides that enriched for positively charged peptides. Also included are a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Characterizing phosphorylated polypeptides in a sample comprises digesting the polypeptides with a protease thus generating test peptides, and collecting a fraction of test peptides that enriched for positively
                                                                                                                                                                            Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 16; Page 69; 123pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2005-031720/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            phosphoprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human phosphorylated peptide from phophoprotein #305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HARD ) HARVARD COLLEGE
                                              12 GEADAMSLDGG
        GEGEAASADDG
                                                                                                                                                                            $
                                                                                        Conservative
                                                                                                              26.1%;
                                                22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                        1;
                                                                                                            Score 30; DB 9;
Pred. No. 2e+03;
                                                                                        Mismatches
                                                                                        4
                                                                                                                                  Length 23;
                                                                                        Indels
                                                                                        0,
                                                                                        Gaps
                                                                                        0
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RESULT 8
AEF54127
ID AEF5
XX AEF5
AC AEF5
XC AEF5
XX CAEF5
XX AEF5
XX AEF7
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                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                 Best Local
                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel isolated polypeptide comprising or consisting of two or more C35 peptide epicopes. The polypeptide of the invention demonstrates cytostatic activity and may be useful for the formulation of an immunogenic composition, such as a vaccine, to induce antibodies and cell-mediated immunity against target cells such as tumour cells. Furthermore, the polypeptide and its analogues may be useful as prognostic markets for carcinoma, such as human breast or bladder carcinoma carcinoma. The current sequence is that of human breast/bladder carcinoma
             therapeutic; prophylaxis; neisseria meningitidis haemophilus influenzae infection; antibacterial;
                                                          N. meningitidis 2996 adhesin NadA HR2a extended SEQ ID NO:30.
                                                                                         23-MAR-2006
                                                                                                                                                AEF54127 standard; peptide; 23
                                                                                                                                                                                                                                                                                                                                          Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 148; 626pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel C35 polypeptide useful for formulation of immunogenic composition to induce antibodies and cell-mediated immunity against tumor cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zauderer M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUN-2002; 2002US-0386738P
11-DEC-2002; 2002US-0432241P
23-APR-2003; 2003US-0464650P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human carcinoma-related C35 epitope / gp100 Pme117 polytope peptide 751.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK49860 standard; peptide; 23 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-062349/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUN-2003; 2003WO-US018252.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003104428-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYRP ) UNIV ROCHESTER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (VACC-) VACCINEX INC
                                                                                                                                                                                                                                                                                                                                                                    polytope polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          epitope; cytostatic; vaccine; tumour; breast; bladder carcinoma;
an; polytope; gp100 Pme117.
                                                                                                                                                                                                                                                      5 SIAKIMNGEADAMSLDG 21
                                                                                                                                                                                                                                                                                 Similarity 6; Conserv
                                                                                                                                                                                                                        TLEKITNSRPPCVLLDG
                                                                                                                                                                                                                                                                                                                                                                       The current sequence is that of human breast/bladder pe polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                            AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Evans EE,
                                                                                         entry)
                                                                                                                                                                                                                                                                                               25.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Borrello MA
                                                                                                                                                                                                                                                                                   Score 29; DB:
Pred. No. 2.8e
3; Mismatches
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                                                                                                                                                                                                                                                                              DB 8; L...
2.8e+03;
                                                                                                                                                                                                                                                                                                             Length 23;
                              infection;
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                                                                                                                                                                                                                                                                                   Gaps
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ARESULT 9
AAG63463
ID AAG6
XX AAG6
XX AAG6
XX AAG6
XX Haem
XX Haem
XX Haem
XX Haem
XX Haem
XX ABOO
XX Synt
XX Synt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a compound that can bind to the heptad repeat sequences HR1 and/or HR2 of the NadA and HadA adhesin on the surface of a meningococcus and haemophilus, respectively, thus inhibiting the ability of the meningococcus and haemophilus either to infect a host organism or to spread an existing infection. The compound is an oligopeptide or polypeptide, or a peptidomimetic compound of the oligopeptide, with anticulated and or anti-haemophilus activity. The invention also includes a pharmaceutical composition comprising the compound and a carrier; a method for treating a patient suffering from a meningococcal or haemophilus infection by administering to the patient the pharmaceutical composition; mutant NadA and HadA proteins, where the mutant protein lacks one or more of the HR1, HR2, or fusion sequences; a nucleic acid encoding the mutant protein; and a bacterium, which expresses the nucleic acid. The compound is useful as a medicament, for the manufacture of a medicament, and for treating a patient for meningococcal or haemophilus infection. This sequence is an extended Neisseria meningitidis 2996 adhesin NadA heptad repeat sequence HR2a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                         Haematopoiesis; haematopoietic stem cell; cell differentiation; haematopoiesis proliferation; myoproliferative syndrome; cytoprotection; apoptosis; medullar expansion; inflammation; anglogenesis; myelofibrosis; thrombocytemia; Vasquez polyglobulia; chronic leukocytic leukemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compound that binds to heptad repeat sequences HR1 or HR2 of NadA and HadA adhesin on the surface of meningococcus and haemophilus, useful for treating meningococcal or haemophilus infection.
                    28-JUN-2001
                                                                                         Synthetic
                                                                                                                                                                                                                                    Amino acid sequence of a peptide which is used in therapy.
                                                                                                                                                                                                                                                                             15-OCT-2001
                                                                                                                                                                                                                                                                                                                                                 AAG63463 standard; peptide; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Masignani V;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUL-2005; 2005WO-IB002320
                                                         WO200146218-A2
                                                                                                                             antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIR ) CHIRON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2006006074-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neisseria meningitidis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTEDSIAKIMN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                             TNKDNIAKKAN 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 28; DB 10;
Pred. No. 4.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 23;
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RESULT 10
AAG63471
ID AAG63
XX
AC AAG62
XX
DE Amin
XX
Heem
KW haem
KW apog
KW haem
KW apog
KW apog
KW apog
KW apog
KW apog
KW bacm
KW apog
KW apog
KW bacm
KW 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG63471 standard; peptide; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-DEC-2000; 2000WO-FR003532.
                                                                                    (VAIS-) INST VAISSEAUX & DU SANG
                                                                                                                                                                22-DEC-1999;
                                                                                                                                                                                                                                        14-DEC-2000; 2000WO-FR003532
                                                                                                                                                                                                                                                                                                                    28-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                WO200146218-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amino acid sequence of a peptide which is used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              subfamily
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  French.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 27; DB 4;
Pred. No. 6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in therapy.
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Alemany M,

Caen JP,

Han ZC;

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RESULT 11
AAG63466
ID AAG63
XX AAG63
AC AAG63
XX Amino
XX Haema
KW Haema
KW Haema
KW Haema
KW Apopt
KW throm
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Best Local S
Matches 5
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                                                                  New eicosa-, tricosa- or pentacosa- to heptacosa-peptides, useful e.g. for stimulating hematopoietic stem cell proliferation after chemotherapy or radiotherapy or as antiinflammatory agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          haematopoiesis proliferation; myoproliferative syndrome; cytoprotection; apoptosis; medullar expansion; inflammation; angiogenesis; myelofibrosis; thrombocytemia; Vasquez polyglobulia; chronic leukocytic leukemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New eicosa-, tricosa- or pentacosa- to heptacosa-peptides, useful e.g. for stimulating hematopoietic stem cell proliferation after chemotherapy or radiotherapy or as antiinflammatory agents.
                                                                                                                                                                                                                                                                                                                     Alemany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-DEC-2000; 2000WO-FR003532.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
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Pred. No.
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6e+03;
Vasquez polyglobulia
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 15; 33pp;
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Pred. No.
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                                                                                        Query Match 23.5%; Score 27; DB 4;
Best Local Similarity 33.3%; Pred. No. 6e+03;
Matches 6; Conservative 4; Mismatches
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                                                   TEDSIAKIMNGEADAMSL 19
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Search completed: June 13, 2006, 02:04:22 Job time: 196 secs

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Result
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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PU00318 PU0036 D60583 PH1693 S47203 I56028 I59174 D33174 B30988 B4845 D64707 B47843 JH0515 B379 B37843 JH0515 PX0011

ALIGNMENTS

RESULT 1 S54339

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homeodomain protein hrox6 - California red abalone (fragment)
C;Species: Haliotis rufescens (California red abalone)
C;Date: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 05-Oct-2004
C;Accession: S60568
R;Degnan, B.M.; Morse, D.E.
Mol. Marine Biol. Biotechnol. 2, 1-9, 1993
A;Reference number: S60564; MUID:93372986; PMID:7689904
A;Accession: S60564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytochrome P450 CYP2A7AS - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: 27-Cct-1995 #sequence_revision 03-Nov-1995 #text_change 07-May-1999
C;Accession: S54339
R;Ding, S.; Lake, B.G.; Friedberg, T.; Wolf, C.R.
Blochem. J. 306, 161-166, 1995
A;Title: Expression and alternative splicing of the cytochrome P-450 CYP2A7.
A;Reference number: S54338; MUID:95169049; PMID:7864805
A;Accession: S54339
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S60568
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                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-23 <DEG>
A;Residues: 1-23 <DEG>
A;Cross-references: UNIPROT:Q25137; UNIPARC:UPI00007984D; EMBL:X79375; NID:g495116; PID:C;Keywords: DNA binding; homeobox; nucleus; transcription regulation
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A;Residues: 1-23 <DIN>
A;Cross-references: UNIPARC:UPI000017C0D9
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Pred. No. 6.2e+02;
""matches 4;
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Pred. No. 8e+03;
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nomeodomain protei

RESULT

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mannose-1-phosphate guanylyltransferase (EC 2.7.7.13) 37K beta chain - pig N;Alternate names: GDP-mannose pyrophosphorylase 37K beta chain C;Species: Sus scrofa domestica (domestic pig) C;Date: 25-Feb-1994 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004 C;Accession: A47415 R;Szumilo, T.; Drake, R.R.; York, J.L.; Elbein, A.D. J. Biol. Chem. 268, 17943-17950, 1993 A;Title: GDP-mannose pyrophosphorylase. Purification to homogeneity, proper A;Reference number: A47415; MUID:93352609; PMID:7688733 A;Contents: liver A47415; MUID:93352609; PMID:7688733 A;Accession: A47415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Description: generates GDP-mannose and pyrophosphate from mannose-1-phosphate and GTP A;Note: also catalyzes synthesis of GDP-glucose from glucose-1-phosphate (EC 2.7.7.34 ac C;Superfamily: mannose-1-phosphate guanylyltransferase C;Keywords: nucleotidyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPROT:Q9TRF4; UNIPARC:UPI00000872E8
A;NOte: sequence extracted from NCBI backbone (NCBIP:136438)
C;Complex: The enzyme appears to be a heterodimer of alpha and beta C;Function:
A;Molecule type: protein A;Residues: 1-23 <KER> A;Cross-references: UNIPF
                                                         A;Reference number: A54226;
A;Accession: B54226
A;Status: preliminary
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A;Molecule type: mRNA
A;Residues: 1-23 <MCG>
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Am. J. Respir. Cell Mol. Biol. 8, 201-208, 1993
A;Title: Cystic fibrosis gene and protein expression during
A;Reference number: I51922; MUID:93152187; PMID:7678968
A;Accession: I51922
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fimbrial adhesin CF29K - Klebsiella pneumoniae (fragment)
C;Species: Klebsiella pneumoniae
C;Species: Klebsiella pneumoniae
C;Date: 10-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A43872
R;Darfeuille-Michaud, A.; Jallat, C.; Aubel, D.; Sirot, D.; Rich, C.; Sirot, J.; Joly, B
Infect. Immun. 60, 44-55, 1992
A;Title: R-plasmid-encoded adhesive factor in Klebsiella pneumoniae strains responsible
A;Reference number: A43872; MUID:92104695; PMID:1345909
A;Accession: A43872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 7
A43872
                                                                                                                                                            A;Cross-references: UNIPROT:Q9R2G4; UNIPARC:UP100000B3816
A;Experimental source: strain CF504, R plasmid
A;Note: sequence extracted from NCBI backbone (NCBIP:7436)
C;Superfamily: fimbrial adhesin K88
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R;Forst, S.; Waukau, J.; Leisman, G.; Exner, M.; Hancock, R. Mol. Microbiol. 18, 779-789, 1995  
A;Title: Functional and regulatory analysis of the OmpF-like pol A;Reference number: S70779; MUID:96414477; PMID:8817498
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C;Keywords: aminotransferase
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A; Residues: 1-23 < FOR>
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Pred. No. 1
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Pred. No. 1.1e+04;
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backbone (NCBIP:74360)
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human

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R;Maier, R.M.; Neckermann, K.; Igloi, G.L.; Koessel, H. J. Mol. Biol. 251, 614-628, 1995
A;Title: Complete sequence of the maize chloroplast genome: A;Reference number: S58531; MUID:95395841; PMID:7666415
A;Accession: S58624
                                                                                                                             hypothetical protein 23 - maize chloroplast
C;Species: chloroplast Zea mays (maize)
C;Date: 29-Nov-1995 #sequence_revision 19-Jan-1996 #text_change 09-Jul-2004
C;Accession: S58604, S58609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 05-Nov-1999 (;Accession: S47194 R;Plaza, A.; Kono, D.H.; Theofilopoulos, A.N. submitted to the EMBL Data Library, February 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mol. Gen. Genet. 217, 185-194, 1989
A;Title: The complete sequence of the rice
of the cereals.
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A; Accession: S47194
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A; Residues: 1-23 <HIR>
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A;Status: nucleic acid sequence not
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A;Accession: JQ0281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-23 < PLA>
A;Molecule type: DNA
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Best Local
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                    Status: nucleic acid sequence
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Pred. No. 1.6e
2; Mismatches
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                      shown;
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                        translation
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1.6e+04;
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A; Note:
C; Keywor
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C;Keywords: chloroplast
                                                                                                                                                                                        A; Molecule type: protein A; Residues: 1-23 < GOL>
                                                                                                                                                                                                                       A; Reference number: S78411
A; Accession: S78417
                                                                                                                                                                                                                                                                                                          ribosomal protein RL31, mitochondrial [validated] - C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #
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C;Genetics: <GEN2>
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A; Map position: IR(A)
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A;Note: the nucleotide
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                                                                                                                                  Cross-references: UNIPARC:UPI000017C9B5; Note: the protein is designated as mitochondrial ribosomal; Reywords: mitochondrion; protein biosynthesis; ribosome
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6; Conservative
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STYXSLXDDYNXKVELALXXDG
                                                                  Conservative
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                                                                                 16.1%;
27.3%;
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Database,
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Pred. No. 1.9e+04;
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1.6e+04;
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A;Molecule type: DNA; A;Residues: 1-23 <RES; A;Cross-references: UNIPARC:UPI00000008D5; GB:M11892; NID:g191914; PIDN:AAA37225.1; C;Superfamily: alpha-amylase, standard type; alpha-amylase core homology

PID:

A;Reference number: I49489; MUID:86033801; PMID:2414282

A;Status: preliminary; translated

from

GB/EMBL/DDBC

Matches Query Match

Local Similarity les 3; Conserv

Conservative

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Mismatches

Indels

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Gaps

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15.7%;

Score 18; DB 2; Pred. No. 2.3e+04;

Length 23;

R;Gumucio, D.L.; Wiebauer, K.; Dranginis, A.; Samuel J. Biol. Chem. 260, 13483-13489, 1985
A;Title: Evolution of the amylase multigene family:

A.; Samuelson, L.C.; Treisman, L.O.; Caldwell,

express a pancreatic

amy

Accession: 149492

B-1 alpha-amylase - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 31-Dec-2004

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RESULT 12

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photosystem I 8.0K G chain - common tobacco (fragment)
C;Species: Nicotiana tabacum (common tobacco)
C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 17-Mar-1999
C;Accession: PQ0690
R;Obokata, J.; Mikami, K.; Hayashida, N.; Nakamura, M.; Sugiura, M.
Plant Physiol. 102, 1259-1267, 1993
A;Title: Molecular heterogeneity of photosystem I. psaD, psaE, psaF, psaH an A;Reference number: PQ0667; MUID:94105345; PMID:8278548
A;Accession: PQ06690
A;Molecule type: protein
A;Residues: 1-23 <080>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    slow muscle tropomyosin - Atlantic salmon (fragment)
C;Species: Salmo salar (Atlantic salmon)
C;Otate: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 09-May-1997
C;Accession: S66449
R;Heeley, D.H.; Bieger, T.; Waddleton, D.M.; Hong, C.; Jackman, D.M.; McGowan, C.; Dav. Bur. J. Biochem. 232, 226-234, 1995
A;Title: Characterisation of fast, slow and cardiac muscle tropomyosins from salmonid: A;Accession: S66449; MUID:96048051; PMID:7556155
A;Accession: S66449; MUID:96048051; PMID:7556155
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                                                                                                                                                                                                                                                                                              RESULT 15
PQ0690
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A;Residues: 1-23 <HEE>
A;Cross-references: UNIPARC:UPI00001775DE
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Best Local Similarity
Watches 3; Conserve
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A;Gene: mGK-6
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
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100.0%; Pred. No. 2.3e+04;
/ative 0; Mismatches 0;
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C;Superfamily: photosystem I chain V
C;Keywords: chloroplast; photosynthesis; photosystem I; thylakoid
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Best Local Similarity 100.0%; Pred. No. 2.3e+04;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps
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Search completed: June 13, Job time : 41 secs

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07-FEB-2006, entry version 10.
Hypothetical protein (Fragment).
ORFManes=PY04268;
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                                                                                                                                                                                                                                                                                                                                                                                                                               "Genome sequence and comparative analysis of the model rodent parasite plasmodium yoelii.";
Nature 419:512-519(2002).

-1- CAUTION: The sequence shown here is derived from an energy of the sequence shown here is derived from an energy of the sequence shown here is derived from an energy of the sequence should be sh
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                                                                                                                                                                                        EMBL; AABL01001282; EAA16099.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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01-FEB-2006, entry version 12.
Collagen type IV 26 kDa component (Fragment).
Canis familiaris (Dog).
Cutte familiaris (Dog).
Cutte familiaris (Dog).
Canis familiaris (Chordata; Craniata; Vertebrata; Euteleostomi; Canida
                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE. Kodatani H., Mignot E.; Submitted (NOV-1998) to
                                                                                       NUCLEOTIDE SEQUENCE.

MEDILINE=99265967; PubMed=10331940; DOI=10.1006/geno.1999.

Li R., Mignot E., Faraco J., Kadotani H., Cantanese J., Z

Lin X., Hinton L., Ostrander E.A., Patterson D.F., de Jon

"Construction and characterization of an eightfold redund
genomic bacterial artificial chromosome library.";

Genomics 58:9-17(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1999, integrated into 0
01-NOV-1999, sequence version
07-FEB-2006, entry version 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROTEIN SEQUENCE.
Thorner_P.S., Baumal R.,
                                                                                                                                                                                                                                                                                                                                                                                            Name=ADA;
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01-NOV-1996, integrated into UniProtKB/TrEMBL
01-NOV-1996, sequence version 1.
07-FEB-2006, entry version 16.
Glycoprotein C precursor (Fragment).
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24-MAY-2005, sequence version 1.
07-FBB-2006, entry version 3.
Gene for alpha 2 (IV) collagen (exon 7) (Fragment).
Name=gC;
Suid herpesvirus 1 (Pseudorabies virus)
Viruses; dsDNA viruses, no RNA stage; H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=85296379; PubMed=3839908; Kurkinen M., Bernard M.P., Barlow D.P., Chow L.T.; Kurkinen M., Bernard M.P., Barlow D.P., Chow L.T.; "Characterization of 64. 123- and 182-base-pair exons in alpha 2(IV) collagen gene."; Nature 317:177-179(1985).
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Murcidea; Muridae; Murinae; Mus.
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nes 3; Conserv
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RESULT 9
Q9PRS0 GADMO
O9PRS0_GADMO PRELIMINARY;
ID Q9PRS0;
AC Q9PRS0;
DT 01-MAY-2000, integrated into the company of the compa
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Q69391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=95081163; PubMed=7989378;
Tomilo M., Wilkinson K.S., Ryan P.;
"Can a signal sequence become too hydrophobic?";
J. Biol. Chem. 269:32016-32021(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suid herpesvirus 1 (Pseudorabies virus)
Viruses; dsDNA viruses, no RNA stage; H
Alphaherpesvirinae; Varicellovirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=Becker;
MEDLINE=95081163; PubMed=7989378;
Tomilo M., Wilkinson K.S., Ryan P.;
"Can a signal sequence become too hyd:
J. Biol. Chem. 269:32016-32021(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; L36969; AAA79966.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=Becker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10345;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
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NCBI_TaxID=10345;
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7 AA;
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Pred. No. 7.
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7.5e+04;
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integrated into UniProtKB/TrEMBL

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RESULT 10
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P56244;
15-JUL-1998, integrated into UniProtKB/Swiss-Prot.
15-JUL-1998, sequence version 1.
07-FEB-2006, entry version 27.
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                              J. Chem. Res. 138:910-936(1993).

-I- FUNCTION: Antibacterial peptide, that adopts an conformation which can disrupt bacterial membras displays a different antimicrobial specificity.
-I- SUBCELLULAR LOCATION: Secreted protein.
-I- TISSUE SPECIFICITY: Expressed by the skin parote
                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Parotoid gland;
Stone D.J.M., Waugh R.J.,
"Peptides from Australian
Litoria caerula.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Litoria caerulea (Green tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Raae A.J., Flengsrud R., Sletten K.; "Chymotrypsin isoenzymes in Atlantic cod; differences substrate specificity."; Comp. Biochem. Physiol. 112B:393-398(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Paracanthopterygii; Gadiformes; Gadidae; Gadus.
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MEDLINE=96043258; PubMed=7584866; DOI=10.1016/0305-0491(95)00094-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=8049;
                                                                                                                                                                                                                                    Amidation; Amphibian defense peptide; Antibiotic; Antimicrobial;
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Local Similarity hes 5; Conserv
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23 AA;
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2356 MW;
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  Score 21; DB 1; Length 23; Pred. No. 1.1e+05; o; Mismatches 6; Indels
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Pred. No. 7.
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RESULT 12
Q9HCW9 HUM
ID Q9HCW
AC Q9HCW
AC Q9HCW
DT 01-MA
DT 01-MA
DT 01-MA
DT 07-FE
DE Growt
GN Name=
OS Homo
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G6PI CAL
P83780;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JAN-2005, sequence version 1.
07-FEB-2006, entry version 6.
Glucose-6-phosphate isomerase (EC 5.3.1.9)
isomerase) (PGI) (Phosphohexose isomerase)
                               01-MAR-2001, integrated into UniProtKB/TrEMBL 01-MAR-2001, sequence version 1. 07-FEB-2006, entry version 10.
                                                                 Q9HCW9;
                                                                         9 HUMAN
Q9HCW9 HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Candida albicans (Yeast).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=PGI1
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                                                                                                                                                                                                                                                                                CHAIN
                                                                                                                                                                                                                                                                                                               InterPro; IPR001672; G6P_Isomerase.
PROSITE; PS00765; P_GLUCOSE_ISOMERASE_1;
PROSITE; PS00174; P_GLUCOSE_ISOMERASE_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- CATALYTIC ACTIVITY: D-glucose 6-phosphate =
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                                                                                                                                                                                                                                                                                                   Antigen; Direct protein

    -!- SIMILARITY: Belongs to the GPI family.

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                                                                            PRELIMINARY;
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                                                                                                                                                                                                                                                                                                   sequencing; Gluconeogenesis;
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Pred. No.
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isomerase) (PHI)
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Growth hormone receptor

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Homo sapiens (Human)

Name=GHR;

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RESULT 13

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Q25137_F
Q25137;
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"Identification of eight homeobox-containing transcripts expressed during larval development and at metamorphosis in the gastropod mollusc Hallotis rufescens";
Mol. Mar. Biol. Biotechnol. 2:1-9(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
Orlovsky I.V., Sverdlova P.S.,
Submitted (JUL-2000) to the EM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (MAY-1994) to the EMBL/GenBank/DDBJ databases.
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MEDLINE=93372986; PubMed=7689904;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hallotis rufescens (California red abalone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hrox6 protein (Fragment).
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11
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                                                                                                        Score 21; DB 2;
Pred. No. 1.1e+05;
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Pred. No. 1.1e+05;
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Attribution-NoDerivs License
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Q4XCL2 PLACH
ID Q4XCL2 PAC
DT Q5-JUL-2
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Q7RMK7;
15-DEC-2003, integ
15-DEC-2003, seque:
07-FEB-2006, entry
Hypothetical prote
ORFNames=FY02172;
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MEDLINE=22255706; PubMed=12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Koolj T.W., Pertea M., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M., Florens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A., Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
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Q4XCL2;
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Science 307:82-86(2005).
-i- CAUTION: The sequence shown here is derived for EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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PubMed=15637271; DO:
                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE [LARGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=73239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmodium yoelii yoelii.
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /EMC Celerra SIDS3/ptodata/2/iaa/5_COMB.pep:*

2: /EMC Celerra SIDS3/ptodata/2/iaa/6_COMB.pep:*

3: /EMC_Celerra SIDS3/ptodata/2/iaa/7_COMB.pep:*

4: /EMC Celerra SIDS3/ptodata/2/iaa/H_COMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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US-08-09-3-3-83-27
US-08-09-128-450-7
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Sequence 14, Appl
Sequence 25, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 28, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 19, Appl
Sequence 13, Appl
Sequence 52, Appl
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Sequence 34, Appl Sequence 215, App Sequence 215, App		nce 5	Patent No. 5221732 Patent No. 5221732 Patent No. 5254535	Sequence 146, App Sequence 54, Appl Sequence 4, Appli	Sequence 53803, A Sequence 58329, A	Sequence 1, Appli Sequence 19, Appl

ALIGNMENTS

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US-08-568-1478-14
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TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/568
FILING DATE:
FILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 800,952
FILING DATE: 02-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: DIGIGITO, Frank S.
REGISTRATION NUMBER: 31,346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: pept
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APPLICANT: Sekiquchi, Kiyoshi
APPLICANT: Takeda, Katsumichi
TITLE OF INVENTION: DNA FRAGMENT CODING FOR SQUAMOUS CELL
TITLE OF INVENTION: CARCINOMA-ASSOCIATED ANTIGEN
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                    REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 8425
TELECOMMUNICATION INFORMATION:
TELEPHONE: 516-742-4343
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COUNTRY: US
ZIP: 11530
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T: 400 Garden City Plaza
Garden City
E: New York
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US-08-178-477B-17
                                                                                                                                                                                                                                                                                                                                               US-08-178-477B-17
                                                                          GENERAL INFORMATION: APPLICANT: HEMKER,
                                                                                                           Sequence 25, Application US/08860808E Patent No. 6207399
                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17, Application US/08178477B Patent No. 5756343
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/617,910
FILING DATE: 26-NOV-1990
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: CAROL M. GRUPPI
REGISTRATION NUMBER: 37,341
REFERENCE/DOCKET NUMBER: 2026-4103U:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: WU, CA
APPLICANT: WESTWC
            TITLE OF INVENTION:
                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: JEM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: WU, CARL; CLOS, JOACHIM;
APPLICANT: WESTWOOD, J. TIMOTHY; RABINDRAN, SRIDHAR
TITLE OF INVENTION: CELL STRESS
TITLE OF INVENTION: TRANSCRIPTIONAL FACTORS
NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 345 PARK AVENUE CITY: NEW YORK
                                                                                                                                                                                                                                                                            Match 24.3%;
Local Similarity 43.8%;
Les 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/178,477B
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                                                         HEMKER, HENDRIK C
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GODEFRIDUS I
METHODS OF DETERMINING ENDOGENOUS THROMBIN POTENTIAL
(etp) AND THROMBIN SUBSTRATES FOR USE IN SAID METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17:
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                                                                                                                                                                                                                                                                                               Score 28; DB 1;
Pred. No. 7e+02;
                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                              Length 23;
                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                              ; MOLECULE TYPE: protein
US-07-936-163-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-07-936-163-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 23.5%;
Best Local Similarity 41.7%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: HEMKER
CURRENT APPLICATION NUMBER: US/08/860,808E
CURRENT FILING DATE: 1997-09-05
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 25
LENGTH: 23
                                                                            Matches
                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READALE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/936,163
FILING DATE: 27-AUG-1992
CLASSIFICATION: 514
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acide
                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: ZINDRICK, THOMAS D
REGISTRATION NUMBER: 32,185
REFERENCE/DOCKET NUMBER: C-38,424A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 49
CORRESONE ADDRESS:
ADDRESSEE: THOMAS D. ZINI
STREET: 9002 PURDUE ROAD
CITY: INDIANAPOLIS
                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: INSECTICIDAL PROTEINS AND METHOD FOR TITLE OF INVENTION: PLANT PROTECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                                                                                                          LENGTH: 23 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                          Local Similarity
les 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 46268-1189
                                                                                                                                                                                                                                                                                 ELEPHONE:
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                                           17 MSLDGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ន
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                                                                          Conservative
                                                                                                                                                                              linear
            14
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                                                                                                                                                                                                                                                                               636-1869
                                                                                            22.6%;
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                                                                            Score 26; DB Pred. No. 1.5e Pred. No. 1.5e
                                                                              2;
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3; Mismatches
                                                                                            DB 1;
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US-08-093-383-27
; Sequence 27, Application US/08093383
; Patent No. 5489529
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEXX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
TYPE: amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 6; Conserv
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Patent No. 5489529
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FILING DATE: 28-NOV-1990
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/198824
FILING DATE: 05-APR-1988
PRIOR APPLICATION DATA:
APPLICATION UMBER: 06/632361
FILING DATE: 19-JUL-1984
PRIOR APPLICATION DATA:
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MEDIUM TYPE: 5.25 inch, 360 K
COMPUTER: IBM PC comparible
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-D
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                               GENERAL INFORMATION:
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APPLICANT: DeBoer, Herman A.
APPLICANT: Heyneker, Herbert L.
APPLICANT: Seeburg, Peter H.
TITLE OF INVENTION: DNA for Expression of Bovine Growth Hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
07/619827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3562
TELEPAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: DeBoer, Herman A.
APPLICANT: Heyneker, Herbert L.
APPLICANT: Seeburg, Peter H.
TITLE OF INVENTION: DNA for Expression of Bovine Growth Hormone
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Johnston, Sear REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/093,383 FILING DATE: 14-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sean A., Sean A. P35,910
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75.0%;
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0; Mismatches
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 27
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
TENGTH: 23 amino acids
                                                                                                                                                                                                                                                                                                                      Sequence 26, Application US/08808550 Patent No. 5871992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMERS: 07/619827
FILING DATE: 28-NOV-1990
PRIOR APPLICATION DATA:
APPLICATION UNMERS: 07/198824
FILING DATE: 05-APR-1988
PRIOR APPLICATION DATA:
APPLICATION UNMERS: 06/632361
PRILING DATE: 19-JUL-1984
PRIOR APPLICATION DATA:
APPLICATION UNMERS: 06/303687
APPLICATION DATA:
APPLICATION UNMERS: 06/303687
FILING DATE: 18-SEP-1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb fl.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/093,383
FILING DATE: 14-JUL-1993
                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                      APPLICANT: Teebor, George W.
APPLICANT: Hilbert, Timothy P.
TITLE OF INVENTION: MAMMALIAN ENDONUCLEASE III AND
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC USES THEREOF
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                         NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.
ADDRESSEE: David A. Jackson, Continental Plaza, 4th
STREET: 411 Hackensack Ave, Continental Plaza, 4th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Johnston, Sean A.
REGISTRATION NUMBER: P35,910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3562
TELEFAX: 415/952-9881
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ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                               CITY: Hackensack
STATE: New Jersey
                                                                                                                  STREET: 411 Ha
                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 AMSLDGGF 23
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                                                                USA
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75.0%;
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Pred. No. 2.2e+03;
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RESULT 9
US-09-220-731-18
; Sequence 18, Application US/09220731A
; Patent No. 6232088
; GENERAL INFORMATION:
, APPLICANT: Phairson Medical, Inc.
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US-09-128-450-7
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                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/128,450
CURRENT FILING DATE: 1998-08-03
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
LENGTH: 23
TYPE: PRT
ORGANISM: Hamster sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/09128450 Patent No. 6211149
GENERAL INFORMATION:
                                                                                                                                                                                                                Matches
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SOFTMARE: PATENTINE Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/808,550
FILING DATE: 26-FEB-1997
CLASSIFICATION: 435
ATTORNEY/AFPNT
                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Chesebro, Bruce W
APPLICANT: Caughey, Byron W
APPLICANT: Chabry, Jelle
APPLICANT: Priola, Susette
TITLE OF INVENTION: Inhibitors of Formation of Protease Resistant Prion
TITLE OF INVENTION: Protein
FILE REFERENCE: 50121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: pepti HYPOTHETICAL: NO FRAGMENT TYPE: inter ORIGINAL SOURCE: ORGANISM: bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1049-1-001 N
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
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Local Similarity 31.8%;
les 7; Conservativo
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Pred. No. 3.2e+03;
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Pred. No. 3.2e+03;
2; Mismatches 6
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RESULT 11 US-08-981-527A-17

Sequence 17, Application US/08981527A Patent No. 6410262 GENERAL INFORMATION: APPLICANT: Quax, Wilhelmus J.

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US-09-823-494-7
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                                                                                                                                                     US-09-823-494-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/220,731A
CURRENT FILING DATE: 1998-12-24
EARLIER APPLICATION NUMBER: PCT/SE93/00455
EARLIER FILING DATE: 1993-05-21
EARLIER FILING DATE: 1993-05-21
EARLIER APPLICATION NUMBER: 08/338,501
EARLIER APPLICATION NUMBER: 08/385,540
EARLIER APPLICATION NUMBER: 08/486,820
EARLIER FILING DATE: 1995-02-08
EARLIER FILING DATE: 1995-02-08
EARLIER FILING DATE: 1995-02-08
EARLIER APPLICATION NUMBER: 08/466,820
EARLIER APPLICATION NUMBER: 08/600,273
EARLIER APPLICATION NUMBER: 08/600,273
EARLIER APPLICATION NUMBER: 08/600,273
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Best Local S
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CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/128,450
PRIOR FILING DATE: 1998-08-03
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/09823494
Patent No. 6355610
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 18
LENGTH: 23
TYPE: PRT
                                                                          Matches
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                                                                                              Best Local
                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chesebro, Bruce W
APPLICANT: Caughey, Byron W
APPLICANT: Chabry, Joelle
APPLICANT: Priola, Susette
TITLE OF INVENTION: Inhibitors of Formation
TITLE OF INVENTION: Protein
FILE REFERENCE: 50121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Richard L. Franklin
APPLICANT: Yves St. Pierre
TITLE OF INVENTION: Treatment and Prevention
TITLE OF INVENTION: Rejection Reactions
FILE REFERENCE: 314572-101D
                                                                                                                                                                     LENGTH: 23
TYPE: PRT
ORGANISM: Hamster sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10 HSQAHQVSLNSGY 22
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                                                                        l Similarity
7; Conserv
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                                    8 KIMNGEADAMSLDGG 22
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H | | | :: | | KHMAGAAAAGAVVGG
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Pred. No.
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Pred. No. 3.
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                                                                                              3.2e+03;
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                                                                          Gaps
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APPLICANT: Kerkman, Richard
APPLICANT: Broekhuizen, Cornelis P.
TITLE OF INVENTION: No. 6410262el Secretion Fa
TITLE OF INVENTION: of Using It
TITLE OF INVENTION: of Using It
FILE REFERENCE: GCX322-US
CURRENT APPLICATION NUMBER: US/08/981,527A
CURRENT APPLICATION NUMBER: PCT/NL96/00278
PRIOR APPLICATION NUMBER: PCT/NL96/00278
PRIOR FILING DATE: 1997-12-16
PRIOR FILING DATE: 1996-07-05
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 23
TYPE: PRT
CURANTEM: Sulfolobus solfataricus
                                                                                                       RESULT 13
US-09-270-767-60291
US-09-270-767-60291, Application US/09270767; Patent No. 6703491; GENERAL INFORMARTION:
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GENERAL INFORMATION:
APPLICANT: Quax, Wilhelmus J.
APPLICANT: Kerkman, Richard
APPLICANT: Broekhuizen, Cornelis P.
TITLE OF INVENTION: No. 6692951el Secretion Factors for
TITLE OF INVENTION: Gram-Positive Microorganisms Genes Encoding Them and Methods
TITLE OF INVENTION: of Using It
FILE REFERENCE: GCX322-US
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Best Local Similarity
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APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767

CURRENT FILLING DATE: 1999-03-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/147,936B
CURRENT FILING DATE: 2002-05-17
PRIOR APPLICATION NUMBER: US/08/981,527
PRIOR FILING DATE: 1997-12-16
PRIOR APPLICATION NUMBER: PCT/NL96/00278
PRIOR APPLICATION NUMBER: PCT/NL96/00278
PRIOR FILING DATE: 1996-07-05
NUMBER OF SEQ ID NOS: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Sulfolobus solfataricus -08-981-527A-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 23
TYPE: PRT
ORGANISM: Sulfolobus solfataricus
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4; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INVENTION: No. 6410262el Secretion Factors for INVENTION: No. 6410262el Secretion Factors for INVENTION: Gram-Positive Microorganisms Genes Encoding Them and Methods INVENTION: of Using It
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nilarity 57.1%;
Conservative
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Pred. No. 3.2e+03;
3; Mismatches 0
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Pred. No. 3.2e+03;
                                                                                                                                                                                                                                                                                                                                                                  0,
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Query Match
Best Local Similarity
Watches 5; Conservi
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                                                                ; ORGANISM: Sus
US-10-071-214-5
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                                                                                                                       Sequence 5, Application US/10071214

Patent No. 7019194

GENERAL INFORMATION:
APPLICANT: HANSSON, Lennart
APPLICANT: HANSSON, Lennart
APPLICANT: HANSSON, Lennart
APPLICANT: EGELRUD, Torbjorn
TITLE OF INVENTION: SCCE MODIFIED TRANSGENIC MAMMALS AND THEIR USE AS MODELS OF HUMAN
FILE REFERENCE: HANSSON-3A

CURRENT APPLICATION NUMBER: US/10/071,214

CURRENT FILING DATE: 2002-02-11
PRIOR APPLICATION NUMBER: US 60/267,422

PRIOR APPLICATION NUMBER: DK PA 2001 00216

PRIOR APPLICATION NUMBER: DK PA 2001 00216

PRIOR FILING DATE: 2001-02-09

NUMBER OF SEQ ID NOS: 50

SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-270-767-60291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: WOLFF, JON
APPLICANT: SOKOLOFF, ALEXANDER
TITLE OF INVENTION: PROCESS FOR UTILIZING EPITOPES RECOGNIZED BY NATURAL ANTIBODIES
FILE REFERENCE: MIRUS.014.02
CURRENT APPLICATION NUMBER: US/09/559,021
CURRENT FILING DATE: 2000-04-27
NUMBER OF SEQ ID NOS: 120
SOCTWARE: PATENTIN VEXESON 3.1
SEQ ID NO 3
LENGTH: 23
LENGTH: 23
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GENERAL INFORMATION:
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SEQ ID NO 60291
LENGTH: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local (
Matches
                                Query Match
                                                                                               LENGTH: 2:
TYPE: PRT
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ORGANISM: Bacteriophage
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Local Similarity
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               20.9%;
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               Score 24; DB 3;
Pred. No. 3.2e+03;
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Pred. No. 3.2e+03;
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Pred. No. 3.2e+03;
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Db 1 AGIPNSKTNACNGDSG 16

Search completed: June 13, 2006, 02:10:59
Job time : 51 secs

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Minimum
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Listing first 45 summaries
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Perfect score:
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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                            SUMMARIES
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Sequence 2, Appli
Sequence 31, Appl
Sequence 63, Appl
Sequence 63, Appl
Sequence 23, Appl
Sequence 21, Appl
Sequence 375, Appl
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Sequence 375, App
Sequence 10, Appl
Sequence 11, Appl
Sequence 17, Appli
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Sequence 54, Appli
Sequence 60, Appli
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US-10-612-162-2
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LENGTH: 23
TYPE: PRT
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19.1	19.1	19.1	19.1	19.1	19.1		19.1	20.0	20.0	20.0	•	20.0	20.0	20.0	20.0	20.0	20.0
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US-10-372-473-13	US-10-377-714-25	US-10-352-786-1902	US-10-352-786-1894	US-10-352-786-1886	-10-352-786-188	-10-174	-09-971-980	US-11-111-953-519	US-11-136-186-146	US-10-839-525-146	US-10-240-641-26	US-10-372-876-447	US-10-109-274A-158	US-10-231-417-445	US-10-083-259-158	US-10-109-171-146	US-10-277-693A-13
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Seguence
•	25, Appl	1902, Ap	1894, Ap	1886, Ap	1881	93, Appl	35, Appl			146, App		447, App	158, App			146, App	•

ALIGNMENTS

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US-10-862-195-357
US-10-862-195-357

| Sequence 357, Application US/10862195
| Publication No. US20050164324A1
| GENERAL INFORMATION:
| APPLICANT: GYGI, STEVEN P.
| TITLE OF INVENTION: SYSTEMS, METHODS AND KITS FOR CHARACTERIZING PHOSPHOPROTEOMES
| FILE REFERENCE: 58890(70207)
| CURRENT FILING DATE: US/10/862,195
| CURRENT FILING DATE: 2004-06-04
| PRIOR APPLICATION NUMBER: 60/476,010
| PRIOR FILING DATE: 2003-06-04
| NUMBER OF SEQ ID NOS: 2245
| SOFTWARE: Patentin version 3.2
| SEQ ID NO 357
| LENGTH: 23
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APPLICANT: Dade Behring Marburg GmbH

TITLE OF INVENTION: Carbohydrate deficient transferrin (CD;

TITLE OF INVENTION: antibodies, their preparation and use

FILE REFERENCE: 2002/B001

CURRENT APPLICATION NUMBER: US/10/612,162

CURRENT FILING DATE: 2003-07-03

PRIOR APPLICATION NUMBER: 10230550.1

PRIOR APPLICATION NUMBER: 10230550.1

PRIOR APPLICATION NUMBER: 03011334.4

PRIOR FILING DATE: 2003-05-19

NUMBER OF SEQ ID NOS: 4

SOFTMARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/10612162 Publication No. US20040014145A1
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NUMBER OF SEQ ID NOS: 76

SOFTWARE: PatentIn version 3.1

SEQ ID NO 63

LENGTH: 23

TYPE: PRT

ORGANISM: Homo sapiens

US-10-484-568-63
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APPLICANT: Lorens, James
APPLICANT: Pray, Todd
APPLICANT: Bennett, Mark
TITLE OF INVENTION: IN VIVO PRODUCTION OF CYCLIC PEPTIDES FOR INHIBITING
TITLE OF INVENTION: PROTEIN-PROTEIN INTERACTION
FILE REFERENCE: A-71433-1/AMP/CYO
CURRENT APPLICATION NUMBER: US/10/422,536
CURRENT FILING DATE: 2003-04-23
PRIOR APPLICATION NUMBER: US 60/187,130
PRIOR APPLICATION NUMBER: US 09/800,770
PRIOR APPLICATION NUMBER: US 09/800,770
PRIOR FILING DATE: 2001-03-06
PRIOR FILING DATE: 2001-03-06
PRIOR FILING DATE: 2001-03-06
PRIOR FILING DATE: 2002-09-30
NUMBER OF SEQ ID NOS: 168
SOFTWARE: Patentin version 3.2
SEQ ID NO 91
LENGTH: 23
TYPE: PRT
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Sequence 63, Application US/10484568
Publication No. US20040248194A1
GENERAL INFORMATION:
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US-10-422-536-91
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Best Local Similarity
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Query Match
Best Local Similarity
                                                                                                                                                                                                APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Lewis, Claire
APPLICANT: Robinson, Jeffrey
TITLE OF INVENTION: Peptide Screen
FILE REFERENCE: 4-32520A/32360/361
CURRENT APPLICATION NUMBER: US/10/484,568
CURRENT FILING DATE: 2004-01-22
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23.5%;
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Pred. No. 1.7e+03;
Score 27; DB 5;
Pred. No. 5e+03;
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                   Length 23;
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APPLICANT: Schafer, Juergen
APPLICANT: Neumann, Thomas
TITLE OF INVENTION: Method For Characterising Analytes
FILE REFERENCE: 01073/0201939-US0
CURRENT APPLICATION NUMBER: US/10/510,246
CURRENT FILING DATE: 2004-10-01
PRIOR APPLICATION NUMBER: PCT/GB03/01485
PRIOR APPLICATION NUMBER: 02252440.9
PRIOR FILING DATE: 2002-04-04
PRIOR FILING DATE: 2002-04-04
NUMBER OF SEQ ID NOS: 58
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; LOCATION: (1)...(23)
; OTHER INFORMATION: anti-angiogenic peptide
US-10-504-905-69
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CURRENT FILING DATE: 2004-08-18
PRIOR APPLICATION NUMBER: PCT/EP03/01698
PRIOR FILING DATE: 2003-02-19
PRIOR APPLICATION NUMBER: GB 0203882.6
PRIOR FILING DATE: 2002-02-19
PRIOR FILING DATE: 2002-02-19
PRIOR APPLICATION NUMBER: GB 0203883.4
PRIOR APPLICATION NUMBER: GB 0203883.4
PRIOR FILING DATE: 2002-02-19
SEQ ID NO 23
LENGTH: 23
TYPE: PRT
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PRIOR FILING DATE: 2002-07-23
NUMBER OF SEQ ID NOS: 90
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 69
LENGTH: 23
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Publication No. US20060040334A1
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APPLICANT: Garcia-Escheverria, Carlos
APPLICANT: Lewis, Claire
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Publication No
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                                                              SOFTWARE: PatentIn Ver.
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                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Xzillion GmbH & Co. KG APPLICANT: Thompson, Andrew H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Robinson, Jeffrey TITLE OF INVENTION: ORGANIC COMPC FILE REFERENCE: DC/4-32360A/32361
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ORGANISM: Homo sapiens
FEATURE:
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No. US20050118598A1
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Pred. No. 5e+03;
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Query Match
Best Local Similarity
6; Conserve
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US-11-069-642-91
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CURRENT APPLICATION NUMBER: US/11/069,642
CURRENT FILING DATE: 2005-02-28
PRIOR APPLICATION NUMBER: 10/232,758
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2003-04-23
PRIOR FILING DATE: 2003-04-23
PRIOR PRILING DATE: 2003-04-23
PRIOR APPLICATION NUMBER: 09/800,770
PRIOR APPLICATION NUMBER: 09/800,770
PRIOR FILING DATE: 2001-03-06
PRIOR FILING DATE: 2001-03-06
PRIOR FILING DATE: 2000-03-06
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                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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APPLICANT: Arrowood, Michael J.
APPLICANT: MOSS, Dolynn M.
TITLE OF INVENTION: ROCCOMBINANT CRYPTOSPORIDIUM PARVUM
TITLE OF INVENTION: ANTIGEN AND DETECTION OF ANTIBODIE
FILE REFERENCE: 14114.0357U3
CURRENT APPLICATION NUMBER: US/09/734,417
CURRENT FILING DATE: 2000-12-11
PRIOR APPLICATION NUMBER: 60/169,797
PRIOR APPLICATION NUMBER: 60/174,054
PRIOR APPLICATION NUMBER: 60/174,054
                                                                                                                                                                                       APPLICANT: Priest, Jeffrey W. APPLICANT: Lammie, Patrick JAPPLICANT: Kwon, James P. APPLICANT: Arrowood, Michael APPLICANT: Moss, Dolynn M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LORENS, JAMES B.
APPLICANT: PRAY, TODD R.
APPLICANT: KINSELLA, TODD M.
APPLICANT: BENNETT, MARK K.
TITLE OF INVENTION: IN VIVO PRODUCTION OF CYCLIC PEPTIDES FOR TITLE OF INVENTION: INHIBITING PROTEIN-PROTEIN INTERACTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: OTHER INFORMATION: Description of Artificial Sequence:labelled OTHER INFORMATION: peptide
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Local Similarity 50.0%;
nes 5; Conservative
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Pred. No. 5e+03;
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                                                                                                                                             ANTIBODIES THERETO
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CURRENT FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 2001-03-27
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                                                                                                          Sequence 362, Application US/10189437 Publication No. US20030194414A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 535
SOFTWARE: Patentin 2.1
SEQ ID NO 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: BOGOCH,
APPLICANT: BOGOCH,
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 23
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Publication No. US20030180328A1
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Matches 5; Conserv
            APPLICANT: BOGOCH, SAMUEL
APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
FILE REFERENCE: 09425/46905
CURRENT APPLICATION NUMBER: US/10/189,437
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVERTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS TITLE OF INVERTION: AND IN INFLUENZA EPIDEMICS FILE REFERENCE: 09425-46904
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Plasmodium falciparum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: VARIANT
LOCATION: 3
OTHER INFORMATION: Xaa = g or s
OTHER INFORMATION: Description of Artificial Sequence:/note
OTHER INFORMATION: synthetic construct
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2002-07-08
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s; Pred. No. 7.1
0; Mismatches
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Pred. No. 7.1e+03;
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. 7.1e+03;
ches 0;
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Sequence 91, Application US/10174410
Publication No. US20030073134A1
GENERAL INFORMATION:
APPLICANT: Louie, Gordon V.
APPLICANT: Buchanan, Sean Grant
APPLICANT: Gajiwala, Ketan S.
APPLICANT: Gajiwala, Ketan S.
APPLICANT: Sauder J. Michael
TITLE OF INVENTION: CRYSTALS AND STRUCTURES OF
TITLE OF INVENTION: 2C-METHLY-D-ERVTHRIFOL, 2,4-1
FILE REFERENCE: 524982000300
CUURRENT APPLICATION NUMBER: US/10/174,410
CUERENT APPLICATION NUMBER: US/10/174,410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/860,050
CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: 60/51,686
PRIOR FILING DATE: 2003-12-23
PRIOR APPLICATION NUMBER: 60/504,958
PRIOR FILING DATE: 2003-09-23
PRIOR PPLICATION NUMBER: 60/476,186
PRIOR PPLICATION NUMBER: 60/476,186
PRIOR PRILING DATE: 2003-06-06
NUMBER OF SEQ ID NOS: 875
SOPTWARE: PATENTIN VERBION 3.2
TYPE: PRT
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PRIOR FILING DATE: 2002-03-26
PRIOR PELICATION NUMBER: 09/984,057
PRIOR FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 729
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Best Local S
Matches 5
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SEQ ID NO 362
LENGTH: 23
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
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Pred. No. 7.1e+03,
4; Mismatches
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ches 6;
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                                                                   2,4-CYCLODIPHOSPHATE SYNTHASE MECPS
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; LENGTH: 23
; TYPE: PRT
; ORGANISM: B. I
US-10-174-410-91
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-400-083-10
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ORGANISM: Hamster sp. US-09-823-494-7
                                NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.
SEQ ID NO 7
LENGTH: 23
TYPE: PRT
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Sequence 10, Application US/10400083
Publication No. US20040186056A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/400,083
CURRENT FILING DATE: 2003-03-20
PRIOR APPLICATION NUMBER: US 10/116,866
PRIOR FILING DATE: 2002-04-05
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/09823494
Publication No. US20010041790A1
GENERAL INFORMATION:
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Matches 5; Conservative
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NUMBER OF SEQ ID NOS: 336
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 91
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APPLICANT: Caughey, Byron W
APPLICANT: Chabry, Joelle
APPLICANT: Priola, Susette
TITLE OF INVENTION: Inhibitors of Formation
TITLE OF INVENTION: Protein
FILE REFERENCE: 50121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ruoslahti, Erkki
APPLICANT: Porkka, Kimmo
APPLICANT: Christian, Sven
APPLICANT: Christian, Sven
TITLE OF INVENTION: HMGNZ Peptides and Related Molecules
TITLE OF INVENTION: that Selectively Home to Tumor Blood Vessels and Tumor Cells
FILE REFERENCE: P-LJ 5662
                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/823,494
CURRENT FILING DATE: 2001-03-30
                                                                                                                              PRIOR APPLICATION NUMBER: 09/128,450 PRIOR FILING DATE: 1998-08-03
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Pred. No.
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Pred. No. 1e+04;
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Sequence 5, Application US/10071214

Publication No. US20030066099A1

GENERAL INFORMATION:
APPLICANT: HANGSON, Lennart
APPLICANT: EGELRUD, Torbjorn

TITLE OF INVENTION: SCCE MODIFIED TRANSGENIC MAMMALS AND THEIR USE AS MODELS OF HUMAN
FILE REFERENCE: HANGSON-3A
CURRENT APPLICATION NUMBER: US/10/071,214
CURRENT FILING DATE: 2002-02-11
PRIOR APPLICATION NUMBER: US 60/267,422
PRIOR FILING DATE: 2001-02-09
PRIOR FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 50
SOFTWARE: Patentin version 3.1
SEQ ID NO 5
LENGTH: 23
TYPE: PRT
CORGANISM: Sus scrofa
US-10-071-214-5
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Search completed: June 13, 2006, 02:24:34 Job time: 180 secs
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5 KHWAGAAAAGAVVGG 19
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Result
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Maximum Match
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Sequence:
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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17.106 Million cell updates/sec
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US-10-787-734-136
US-10-787-734-136
US-10-787-734-139
US-11-25-276-145-457
US-10-731-899-21
US-10-731-899-21
US-10-51-465-53
US-11-169-140-26
US-11-169-140-26
US-11-151-429-31
US-11-219-563-41
US-11-219-563-43
US-11-219-563-43
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                                 13, Appl
11, Appl
114, Appl
124, App
127, Appl
457, Appl
457, Appl
53, Appl
53, Appl
53, Appl
541, Appl
54
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	2601, A	15, Appl		_	644, App			11, App	•		110, Apr				$\overline{}$			12, App	•	-

ALIGNMENTS

US-11-251-465-55
US-11-251-465-55
; Sequence 55, Application US/11251465
; Publication No. US20060094061A1
; GENERAL INFORMATION:

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RESULT 2
US-11-263-057-13
US-11-263-057-13
Sequence 13, Application US/11263057
Publication No. US20060094103A1
GENERAL INFORMATION:
APPLICANT: Chou, Ku-Chen
APPLICANT: Howe, W. Jeffery
TITLE OF INVENTION: Modified BACE
FILE REFERENCE: MBHB 01-1766-A
CURRENT APPLICATION NUMBER: US/11/263,057
CURRENT FILING DATE: 2005-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/619,384
PRIOR FILING DATE: 2004-10-15
NUMBER OF SEQ ID NOS: 880
SOFTWARE: PatentIn version 3.3
SEQ ID NO 55
LENGTH: 23
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Protein domain
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Best Local
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APPLICANT: Klaassen, Hubertus
ITILE OF INVENTION: Molecular Targets And Compounds,
TITLE OF INVENTION: Same, Useful In The Treatment O
TITLE OF INVENTION: Inflammatory Diseases
FILE REFERENCE: P30,172-A USA
CURRENT APPLICATION NUMBER: US/11/251,465
CURRENT FILING DATE: 2005-10-14
CURRENT FILING DATE: 2005-10-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Brys, Reginald APPLICANT: Vandeghinste, Nick
PRIOR APPLICATION NUMBER: US/10/372,473
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Pred. No. 6e+02;
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In The Treatment Of Joint Degenerative And
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US-10-787-734-11
; Sequence 11, Application US/10787734
; Publication No. US20060115489A1
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                                                                                                                                                                                                                                                                    US-10-787-734-136
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; ORGANISM: Influenza A virus
US-10-787-734-11
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CURRENT APPLICATION NUMBER: US/10/787,734
CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/US03/05196
PRIOR FILING DATE: 2003-02-21
PRIOR FILING DATE: 2002-10-21
PRIOR FILING DATE: 2002-10-21
PRIOR PLICATION NUMBER: US 10/274,616
PRIOR PLICATION NUMBER: US 10/082,014
PRIOR APPLICATION NUMBER: US 10/082,014
PRIOR FILING DATE: 2002-02-21
PRIOR FILING DATE: 2002-02-21
PRIOR FILING DATE: 2002-02-31
PRIOR FILING DATE: 2001-08-15
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; NAME/KEY: MISC FEATURE
; OTHER INFORMATION: Synthetic peptide substrate with a P33K-BACE cleavage site.
US-11-263-057-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.2 SEQ ID NO 11
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NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn version 3.2
SEQ ID NO 13
LENGTH: 23
                                                                                                                                                                                                   Sequence 136, Application No. US200 GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: BIRKETT, ASHLEY J.
APPLICANT: BIERS, WALTER
TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE
FILE REFERENCE: 4564/91531
CURRENT APPLICATION NUMBER: US/10/787,734
CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/US03/05196
PRIOR FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: US 10/274,616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE
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 3; Mismatches

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Pred. No. 3.5e+03;
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Pred. No. 1.2e+03;
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PRIOR FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: US 10/082,014
PRIOR ETLING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: US 10/080,299
PRIOR FILING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: US 09/930,915
PRIOR FILING DATE: 2001-08-15
PRIOR FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 140
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 136
LENGTH: 23
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US-11-252-276-124
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Best Local Similarity
Matches 3; Conserve
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US-10-787-734-139
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PRIOR FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: US 10/274,616
PRIOR FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: US 10/082,014
PRIOR FILING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: US 10/080,299
PRIOR FILING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: US 09/930,915
PRIOR APPLICATION NUMBER: US 09/930,915
PRIOR FILING DATE: 2001-08-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 60.0
"...rches 3; Conservative
                                                     Sequence 124, Application US/11252276
Publication No. US20060094868A1
GENERAL INFORMATION:
APPLICANT: Giuliano, Kenneth A.
APPLICANT: Bright, Gary
APPLICANT: Olson, Keith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.2 SEQ ID NO 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: BIRKETT, ASHLEY J.
APPLICANT: FIERS, WALTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 139, Appublication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE FILE REFERENCE: 4564/91531
CURRENT APPLICATION NUMBER: US/10/787,734
CURRENT FILING DATE: 2004-02-26
APPLICANT: Burroughs-Tencza, Sarah
TITLE OF INVENTION: A System for Cell Based Screening
FILE REFERENCE: 97-022-K2-CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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                                                                                                                                                                                                                                                                                                13 EADAMSLDG 21
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Pred. No. 3.5e+03;
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Pred. No. 3.5e+03
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CURRENT APPLICATION NUMBER: US/11/252,276
CURRENT FILING DATE: 2005-10-17

APPLICATION NUMBER: 09/713,572 FILING DATE: 2000-11-15

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OTHER INFORMATION: Cytoplasm/cytoskeleton target seque
US-11-252-276-124
                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,935
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,376
PRIOR APPLICATION NUMBER: 60/424,376
PRIOR APPLICATION NUMBER: 60/424,376
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR PILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR PILING DATE: 2002-11-06
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PRIOR FILLING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 09/398,965
PRIOR FILLING DATE: 1999-09-17
PRIOR APPLICATION NUMBER: 09/031,271
PRIOR FILLING DATE: 1998-02-27
PRIOR APPLICATION NUMBER: 08/810,983
PRIOR FILLING DATE: 1997-02-27
PRIOR FILLING DATE: 1997-02-27
PRIOR APPLICATION NUMBER: 60/136,078
PRIOR APPLICATION NUMBER: 60/136,078
PRIOR APPLICATION NUMBER: 60/136,078
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SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 124
LENGTH: 23
Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 844

SOFTWARE: PatentIn Ver. 3.3

SEQ ID NO 457
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Best Local Similarity
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APPLICANT: DHARAMSI, AKI
APPLICANT: VEDADI, MASQUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
FILE REFERENCE: IPT-330.01
                                                                                                             PRIOR APPLICATION NUMBER: 60/424,362
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,373
PRIOR FILING DATE: 2002-11-06
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CURRENT FILING DATE: 2005-05-05
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Pred. No. 4.9e+03;
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CURRENT APPLICATION NUMBER: US/10/490,949
CURRENT FILING DATE: 2004-03-25
PRIOR APPLICATION NUMBER: PCT/US02/30578
PRIOR FILING DATE: 2002-09-25
PRIOR APPLICATION NUMBER: US 60/339,284
PRIOR FILING DATE: 2001-12-11
PRIOR APPLICATION NUMBER: US 60/325,499
PRIOR PILING DATE: 2001-09-25
NUMBER OF SEO ID NOS: 59
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 36
LENGTH: 23
TYPE: PAT
ORGANISM: Homo Bapiens
US-10-490-949-36
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; Sequence 36, Application US/10490949
; Publication No. US20060093574A1
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SOFTWARE: Patentin version 3.3
SEQ ID NO 21
LENGTH: 23
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Best Local (
Best Local Similarity
                   Query Match
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APPLICANT: ALBANI, Salvotore
APPLICANT: MARTINI, Alberto
TITLE OF INVENTION: METHODS FOR EPITOPE-SPECIFIC AND CYTOKINE/ANTICYTOKINE
TITLE OF INVENTION: COMBINATION IMMUNOTHERAPIES FOR MODULATION OF PATHOGEN
TITLE OF INVENTION: IMMUNE RESPONSES IN IMMUNE MEDIATED DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: CHIMERIC PEPTIDES AS 1
TITLE OF INVENTION: FOR IMMUNIZATION USIT
FILE REFERENCE: 20555/1203433-US1
CURRENT APPLICATION NUMBER: US/09/731,899
CURRENT FILING DATE: 2000-12-08
PRIOR APPLICATION NUMBER: 60/169,687
PRIOR FILING DATE: 1999-12-08
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les 3; Conserv
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42.9%;
14.8%;
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 Score 17;
Pred. No.
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Pred. No. 4.9e+03;
3; Mismatches 7;
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Pred. No. 6.9e+03;
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 DB 6; I
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NUMBER OF SEQ ID NOS: 880
SOFTWARE: Patentin version 3.3
SEQ ID NO 53
LENGTH: 23
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-11-169-140-26
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                                                                   APPLICANT: CHAPIN, Steven
TITLE OF INVENTION: METHODS FOR THE TRANSPORT OF BIOLOGICALLY ACTIVE AGENTS ACROSS CE
TITLE OF INVENTION: METHODS FOR THE TRANSPORT OF BIOLOGICALLY ACTIVE AGENTS ACROSS CE
TITLE OF INVENTION: MARIERS
FILE REFERENCE: 11474-037-999
CURRENT APPLICATION NUMBER: US/11/169,140
CURRENT FILING DATE: 2005-06-27
PRIOR APPLICATION NUMBER: US 60/267,601
PRIOR FILING DATE: 2001-02-09
PRIOR FILING DATE: 2000-11-14
PRIOR APPLICATION NUMBER: US 60/248,819
PRIOR APPLICATION NUMBER: US 60/248,478
PRIOR APPLICATION NUMBER: US 60/248,478
PRIOR APPLICATION NUMBER: US 60/237,929
PRIOR APPLICATION NUMBER: 09/949,039
PRIOR FILING DATE: 2001-09-06
PRIOR FILING DATE: 2001-09-06
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APPLICANT:
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TITLE OF INVENTION: Molecular Targets And Compounds, And Methods To Identify
TITLE OF INVENTION: Same, Useful In The Treatment Of Joint Degenerative And
TITLE OF INVENTION: Inflammatory Diseases
FILE REFERENCE: P30,172-A USA
CURRENT FAPLICATION NUMBER: US/11/251,465
CURRENT APPLICATION NUMBER: US/11/251,465
CURRENT FILING DATE: 2004-10-14
PRIOR APPLICATION NUMBER: 60/619,384
PRIOR FILING DATE: 2004-10-15
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APPLICANT: Vando
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ARIZEKE PHARMACEUTICALS, INC. APPLICANT: HOUSTON, Lou, L. APPLICANT: SHERIDAN, Philip, J.
                   PRIOR APPLICATION NUMBER: 09/969,748 PRIOR FILING DATE: 2001-10-02 NUMBER OF SEQ ID NOS: 143
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1es 5; Conservation
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GLYNN, Jacqueline, M.
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Pred. No. 6.9e+03;
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                                                                                                                                                                                                                                 RESULT 13
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PRIOR FILLING DATE: 2000-12-09
PRIOR PELICATION NUMBER: PCT/US99/01621
PRIOR FILLING DATE: 1999-01-27
PRIOR APPLICATION NUMBER: US 60/073,160
PRIOR FILLING DATE: 1998-01-30
PRIOR PELICATION NUMBER: US 60/073,159
PRIOR APPLICATION NUMBER: US 60/073,159
PRIOR APPLICATION NUMBER: US 60/073,165
PRIOR APPLICATION NUMBER: US 60/073,165
                                                                                                                                Sequence 30, Application US/10515429 Publication No. US20060105387A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 291
LENGTH: 23
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Best Local
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LENGTH: 23
                APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/949,925
CURRENT FILING DATE: 2001-09-12
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CURRENT APPLICATION NUMBER: US/10/515,429
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PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: US 60/073,162
PRIOR FILING DATE: 1998-01-30
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 1998-01-30
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5. US20060099575A9
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Score 17; DB 7;
Pred. No. 6.9e+03;
1; Mismatches 1
      Score 16; DB 1;
Pred. No. 9.7e+03;
                 Length 23
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Indels
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Gaps
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CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779

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APPLICANT: Bander. Neil
APPLICANT: Bander. Neil
TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC
TITLE OF INVENTION: MEMBRANE ANTIBODIES
FILE REFERENCE: 13651.001 (BZL-001)
CURRENT APPLICATION NUMBER: US/11/219,563
CURRENT FILING DATE: 2005-09-02
PRIOR APPLICATION NUMBER: PCT/US04/06586
PRIOR APPLICATION NUMBER: US 10/379,838
PRIOR APPLICATION NUMBER: US 10/379,838
PRIOR APPLICATION NUMBER: 10/449,379
RRIOR APPLICATION NUMBER: 10/449,379
RRIOR FILING DATE: 2003-05-30
NUMBER OF SEQ ID NO 41
SOFTMARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 41
LENGTH: 23
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US-11-219-563-41
; Sequence 41, Application US/11219563
; Publication No. US20060088539A1
; Publication No. US20060088539A1
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PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEO ID NOS: 75
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 30
PRIOR HILLING DATE: 2003-07-09
NUMBER OF SEO ID NOS: 75
SOFTWARE: PATENTIN VER. 2.1
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US-11-259-950-21
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                                                               Sequence 21, Application US/11259950 Publication No. US20060088524A1 GENERAL INFORMATION:
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Best Local Similarity 20.0%;
Matches 2; Conservative
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Best Local Similarity 28.6%;
                                                                                                                                                                                                                                                                                                                     Matches
APPLICANT: Morrissey, James
APPLICANT: Pureza, Vincent
APPLICANT: Sligar, Stephen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: EPO mimetic OTHER INFORMATION: peptide
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Pureza, Vincent
Sligar, Stephen G.
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Pred. No. 9.7e+03;
2; Mismatches E
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Pred. No. 9.7e+03;
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Search completed: June 13, 2006, 02:24:56 Job time : 18 secs
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; OTHER INFORMATION: HisTEV sequence
US-11-259-950-21
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CURRENT FILING DATE: 2005-10-27
PRIOR APPLICATION NUMBER: US 60/622,737
PRIOR FILING DATE: 2004-10-7
PRIOR PRIOR DATE: 2004-01-13
PRIOR PPLICATION NUMBER: US 60/536,281
PRIOR PPLICATION NUMBER: US 10/465,789
PRIOR PILING DATE: 2003-06-18
PRIOR PPLICATION NUMBER: US 09/990,087
PRIOR PILING DATE: 2001-11-20
PRIOR PRIOR DATE: 2001-11-20
PRIOR APPLICATION NUMBER: US 60/252,233
PRIOR PILING DATE: 2000-11-20
PRIOR PILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 11/033,489
PRIOR PILING DATE: 2005-01-11
                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 96
SOFTWARE: PatentIn version 3.3
SEQ ID NO 21
LENGTH: 23
                                                                                                                                                                                 Query Match 13.9%;
Best Local Similarity 75.0%;
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Tissue Factor Compositions and Methods FILE REFERENCE: 46-04
                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial
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                                                                                    TTEN 18
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Pred. No. 9.7e+03;
1; Mismatches 0
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Result
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Maximum DB
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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   protein search, using sw
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Match
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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geneseqp2000s:*
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ABM83778
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ADY54154
AAB84371
ABR82321
ABM83782
ABM83781
ABM83779
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Aee39171 Human ser
Aee39172 Human fer
Adh83932 Human tra
Adk15881 Human tra
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Ady54154 Amino aci
Aab84371 Amino aci
Abr82321 Human dia
Abm83782 Human dia
Abm83781 Human dia
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Adm83778 Human dia
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Adh189360 Human tra
Adh19361 Human tra
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Add45282	Abp72819	Aay50717	Aaw54354	Aar66492	Aar12499	Aee47501	Ady57103	Ady57102	Ady54149	Adt55066	Adu00630	Adu17468	Adu17611	Adt98423	Adt36538	Adt99248	Adt78149	Adt78912	Adu07204	Adt55264	Adu17885
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ALIGNMENTS

RESULT 1 ADG46077

ADG46077;

ADG46077 standard; peptide;

15 B

Human CDT peptide fragment 25-MAR-2004 (first entry)

#3

19-MAY-2003; 2003EP-00011334. EP1378521-A1. antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. 05-JUL-2002; 2002DE-01030550. 07-JAN-2004. Homo sapiens. (DADE-) DADE BEHRING MARBURG GMBH.

Althaus H;

WPI; 2004-073743/08.

New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization.

Claim 4; SEQ ID NO 3; 21pp; German.

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSMGGKEDLIWELL, TTEDSIAKIMNGEADAMSLDGGF, SKLSMGSGLNLSEPN and YEKYLGEEYVKAV (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for

ABM83783 AAU02938 ABM83780 AAU02937

ADK15869 ADH89360 ABP72820

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This invention describes a novel antibody that, in aqueous solution, CC binds selectively to carbohydrate-deficient transferrin (CDT) without chaving to bind CDT to a solid phase. The invention also describes an CC antibody that binds selectively to CDT at regions containing the CC sequences VVARSMGGKEDLINELL, TTEDSIAKINMCBADAVSLDGGF, SKLSMGSGLINESEN and CC YEKYLGEEYVKAV (ADG46075-ADG46078) and an immunoassay for detecting CDT cusing the antibodies of the invention. The antibodies of the invention of are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The CC are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The CC are monoclonal, or a fragment, then generation of hybridomas by standard cC transferrin, or a fragment, then generation of hybridomas by standard for fusion of spleen and myeloma cells. Hybridomas are then selected for production of spleen and myeloma cells. Hybridomas are then selected for CC phase. The antibodies are useful for diagnosis of alcoholism, CDT, which CC alcoholics. The antibodies allow direct detection of CDT in solution, CC alcoholics. The antibodies and/or Asn 611, is present only in CC alcoholics. The antibodies and of crown for a solid phase (as required when using known antibodies). ADG46075-ADG46094 represent CDT peptide
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Matches 15; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating
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non-glycosylated transferrin; alcoholism; alcoholic;
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The invention relates to identifying glycopolypeptides in a serum or CC plasma sample comprising immobilizing derivatized sample CC glycopolypeptides to a solid support, releasing the sample glycopeptide (CC glycopeptide fragments that standard peptides, and identifying released sample CC glycopeptide fragments that correspond to standard peptides added by mass spectroscopy. Also included are a method for identifying one or more CC diagnostic markers for a disease, a composition comprising peptides containing the glycosylation sites (AEE35897-AEE39378, where the peptides containing the glycosylation sites (AEE35897-AEE39378, where the peptides containing the same cleavage reagent) and a kit comprising peptides containing the glycopolypeptides (AEE35897-AEE39378). The methods are useful for clearitying glycopolypeptides in a serum or plasma sample. The methods can also be used to identify and/or containing to the glycopolypeptides in a serum or plasma sample of prognostic and containing to the detection of prognostic and contained containing the state of glycopolypeptides in a laso be used to identify and/or containing to the detection of prognostic and contained the state of glycopolypeptides and true drug efficacy, drug dosing, and/or containing the state of glycopolylation of protein sbased on the concurrent contained to protein abundance measurement of protein glycosylation on the same sample. The method allows fast throughput and simplicity. It can be readily adapted for high throughput analysis of samples, which can be gamples. The method allows fast throughput and simplicity. It can be readily adapted for the facilitate the processing of multiple contains and contains also be automated to facilitate the processing of multiple samples. The method allows fast the processing of multiple contains and contains a human glycopoptide that does not
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rheumatoid arthritis; antiarthritic; antirheumatic; psychiatric disorder;
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each correspond to peptide fragments derived by cleavage of polypeptides using the same cleavage reagent) and a kit comprising peptides containing the glycosylation sites (ARE35897-ARE33378). The methods are useful for identifying glycopolypeptides in a serum or plasma sample. The methods can be used for blood serum profilling for the detection of prognostic and diagnostic protein markers. It can also be used to identify and/or validate drug targets and to evaluate drug efficacy, drug dosing, and/or drug toxicity. The methods can also be used for the detection of changes in the state of glycosylation of proteins based on the concurrent application of protein abundance measurement of protein glycosylation on the same sample. The method allows fast throughput and simplicity. It can be readily adapted for high throughput analysis of samples, which can be
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85.7%;
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Pred. No. 0.0027;
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RESULT 5
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Best Local
This invention relates to a novel fusion protein which comprises a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region. The invention may be useful for the development of compounds with cytostatic, antibacterial, virucide, antiparasitic, immunosuppressive or antiarthritic activity. In addition, the sequences disclosed may be useful for gene therapy. The fusion protein is useful for preparing a composition for treating a disease or disease symptom in a patient for example septic shock, endotoxic shock, cachexia syndromes associated with bacterial, viral or parasitic infections, neoplasm, autoimmune disease, arthritis or adverse effects associated with treatment for preventing graft rejection. The present sequence is that of a domain of the human transferrin protein which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           glycosylation fused to a preparing a composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New fusion protein comprising a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-AUG-2001; 2001US-0315745P;
30-NOV-2001; 2001US-0334059P;
30-AUG-2002; 2002US-00231494,
30-AUG-2002; 2002US-0406977P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiparasitic; immunosuppressive; antiarthritic; gene therapy;
septic shock; endotoxic shock; cachexia syndrome; bacterial infecti
viral infection; parasitic infection; neoplasm; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            particularly advantageous for the analysis of clinical specimens. The method can also be automated to facilitate the processing of multiple samples. The present sequence is a human glycopeptide that does not contain an N-linked glycosylation site, suitable for use as a reference peptide in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                             Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sadeghi H,
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antibody variable region; cytostatic; antibacterial; virucide;
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                       82pp;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                          English.
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No. 0.0027;
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RESULT 6
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RESULT 7
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Matches 12
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises a fusion protein that contains a modified transferrin (Tf) protein fused to a therapeutic protein/peptide (e.g. beta-interferon - IFN, glucagons-like peptide - GLP-1, erythropoietin mimetic peptide - EMP1, T-20 and soluble toxin receptor). The fusion protein of the invention is useful for epitope tagging. The present amino acid sequence represents a fragment of the human Tf protein.
           ADY54154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-AUG-2001; 2001US-0315745P.
30-NOV-2001; 2001US-0334059P.
30-AUG-2002; 2002US-00231499.
30-AUG-2002; 2002US-0406977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fusion protein; transferrin; Tf; beta-interferon; beta-IFN; glucagons-like peptide; GLP-1; erythropoietin mimetic peptide; T-20; soluble toxin receptor; epitope tagging; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 42
                                                          ADY54154 standard; peptide; 42 AA
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New fusion protein comprising a modified transferrin (Tf) protein fused to a therapeutic protein or peptide, useful for epitope tagging.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BIOR-) BIOREXIS PHARM CORP.
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                                                                                                                                                                                                                                 KLSMGSGLNLSEPN 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lai C,
                                                                                                                                                                                                                                                                                                                                                                                                           Ŗ,
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                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Tf) protein fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sadeghi H,
                                                                                                                                                                                                                                                                                                                                81.1%;
85.7%;
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                    Score 60; DB 8;
Pred. No. 0.0042;
0; Mismatches
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Pred. No. 0.0042;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Turner
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                                                                                                                                                                                                                                                                                                                                                      Length 42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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nootropic; hepatotropic; virucide; antiinflammatory; immunosuppressive; antiallergic; antimicrobial; neuroleptic; gynecological; anorectic; antiarteriosclerotic; gastrointestinal-Gen; endocrine-Gen; neoplasm; hematological disease; transferrin; erythropoletin peptide mimetic; EPM; EPO mimetic peptide-1; EMP-1; multiple sclerosis; brain tumor; cancer; hepatitis; anemia; pregnancy; menetrual disorder; rheumatoid arthritis; AIDS; viral disease; metabolic disease; autoimmune disease; inflammatory disease; merabolic disease; neurodegenerative disease; hematopoletic cell disorder; endocrine disorder;
                                                                                                                                                                                                                                                                                                                                                                                                              30-AUG-2004; 2004WO-US027949
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2005021579-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gastrointestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amino acid sequence of C1 subdomain of human transferrin
                                                                                                                                                                                                                                                                                                                                                                                                                                   10-MAR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anti-HIV; hypotensive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease; hypertension; arterial sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cardiovascular-Gen.;
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Novel erythropoietin (EPO) peptide mimetic, having first modification of cysteine residue of EPO mimetic peptides (EMP)-1, to reduces disulfide bond formation, and second modification such that peptide exhibits EMP-1

WPI; 2005-214540/22.

Sadeghi H,

Turner AJ

28-AUG-2003; 2003WO-US026818. 10-MAR-2004; 2004US-0551552P.

(BIOR-) BIOREXIS PHARM CORP

Example 1; SEQ ID NO 8; 158pp; English.

CC (EPM), comprising a modification of at least one cysteine residue of EPO
CC mimetric peptide (EMP)-1 that substantially reduces disulfide bond
CC formation, and a second modification such that the peptide exhibits EMP-1
CC activity. The first modification such that the peptide exhibits EMP-1
CC activity. The first modification such that the peptide exhibits EMP-1
CC activity. The strength of the comprises the deletion or substitution
CC comprises the addition of a linker group that is covalently bonded to the
CC cherminal amino acid or N-terminal amino acid of EMP-1. EPM peptides of
CC the invention are useful for treating or preventing diseases, such as
CC multiple sclerosis, brain tumor, skin cancer, hepatitis B, hepatitis C,
CC anemia, beta-thalassemia, pregnancy or menstrual disease, such as continuous edisease, inflammatory disease, metabolic disease, obesity,
CC autoimmune disease, inflammatory disease, metabolic disease, obesity,
CC disease, systemic microbial infection, cardiovascular disease, psychosis,
CC genetic diseases, neurodegenerative diseases, disorders of hematopoietic
CC cells, diseases, neurodegenerative diseases, disorders of hematopoietic
CC cells, diseases, inflammatory diseases, asthma, or HIV infections,
CC Alzheimer's disease. The presents sequence represents the C1 subdomain of
CC human transferrin A fusion protein, with increased half life, comprising
CC manutal protein and modified transferrin is used in the course of the The specification describes an erythropoietin (EPO) peptide mimetic (EPM), comprising a modification of at least one cysteine residue o invention.

Sequence 42 A

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Matches
      Query Match
Best Local (
12;
      Similarity
Conservative
      81.1%;
      Score 60;
Pred. No.
Mismatches
        0
       0042;
             Length 42
Indels
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N KLSMGSGLNLSEPN 15

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RESULT 8
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ID AAB8
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Matches
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to subburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents human transferrin polypeptide. Transferrin polypeptides and polynucleotides are used to prevent, tre or ameliorate a medical condition in e.g. humans mice, rabbits, goat horses, cats, dogs, chickens or sheep. Disorders which are diagnosed treated include autoimmune diseases e.g. rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; transferrin; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebral ischemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing;
                      ABR82321 standard; protein; 575 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecule encoding a human tranferrin protein is used in preventing, treating or ameliorating a medical condition.
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200146254-A1
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                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                                                                                                                                                                                                        l Similarity
12; Conserv
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                                                                                                                                                                                                                                              KLSMGSGLNLSEPN 15
                                                                                                                                                                                 KLCMGSGLNLCEPN 192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 286-287; 290pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shi Y,
                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                  81.1%;
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                                                                                                                                                                                                                                                                                                        Score 60; DB 4;
Pred. No. 0.042;
0; Mismatches
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                                                                                                                                                                                                                                                                                                     2;
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                                                                                                                                                                                                                                                                                                        Gaps
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cc polynucleotides. The human MEPR polypeptides, polynucleotides and compositions are useful for diagnosing, treating or preventing disorders associated with aberrant expression of MEPR, particularly cell associated with aberrant expression of MEPR, particularly cell cc proliferative disorders (e.g. arteriosclerosis, atherosclerosis, cell providers (e.g. arteriosclerosis, atherosclerosis, cirrhosis, hepatitis, paroxysmal nocturnal hemoglobinuria, polycythemia cc vera, psoriasis, primary thromobocytopenia or cancer), developmental cc disorders (e.g. renal tubular acidosis, anemia or mental retardation), cc neurological disorders (e.g. Alzheimer's disease, parkinson's disease or cepilepsy), autoimmune/inflammatory disorders (e.g. AIDS, allergies, csthmas, autoimmune thyroiditis, contact dermatitis, Crohn's disease, cc diabetes mellitus, glomerulonephritis, Goodpasture's syndrome, gout, Graves' disease, Hashimoto's thyroiditis, irritable bowel syndrome, cmltiple sclerosis, osteoasthritis, osteoporosis, pancreatitis, Reiter's syndrome, rheumatoid arthritis, Sjogren's syndrome, uveitis, or viral, contections, the conding MEPR are useful for creating transgenic animals contections of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEPR; metalloprotein; anti-HIV; antiallergic; antiinflammatory; human; antianemic; antiparkinsonian; nootropic; anticonvulsant; cytostatic; antiarteriosclerotic; antiasthmatic; immunosuppressive; antithyroid; hepatotropic; dermatological; antidiabetic; nephrottropic; antigout; thyromimetic; neuroprotective; osteopathic; antiarthritic; uropathic; antiparasitic; antihelmintic; antipsoriatic; ophthalmological; virucide; antiheumatic; haemostatic; antibacterial; protozoacide; fungicide;
Sequence 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human metalloproteins and polynucleotides, useful for diagnosing, treating or preventing autoimmune or inflammatory disorders (e.g. AIDS, allergy or anemia), multiple sclerosis, osteoarthritis, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JAN-2002; 2002US-0348769P.
18-JAN-2002; 2002US-0350701P.
19-MAR-2002; 2002US-0366059P.
10-MAY-2002; 2002US-0379907P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to novel human metalloproteins (MEPR) and encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 146-148; 153pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hepatitis.
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M, Chien D,
   ΑA;
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Jin P,
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Chawla NK, Yue
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e H, Khare
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KLCMGSGLNLCEPN KLSMGSGLNLSEPN 15

Query Match Best Local Matches

Local Similarity nes 12; Conserv

Conservative

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Mismatches

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Gaps

0

81.1%; 85.7%;

Score 60; DB 6; Pred. No. 0.079;

Length 575; Indels

N

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RESULT 10
RAM83782
ID ABM83782
ID ABM833
XX ABM83
XX ABM83
XX ABM83
XX ABM83
XX Gene
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XX Gene
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XX Gene
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XX 12-SE
XX 12-SE
XX 12-SE
XX I2-SE
XX I
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                                                                                                                                                                                                                                                                                                            The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated CC used muon molecules, e.g. cell proliferative disorders, associated CC autoimmune/inflammatory disorder, developmental disorders, or CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the CC invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly form WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                Best Local
Matches 1
                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV, Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirtor Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gi Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM83782;
                                                                                                                                                                                                                                                                Sequence 609 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-2002;
12-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2003; 2003WO-US028227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004023973-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM83782 standard; protein;
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DB; ACN42434.
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                                                                                                                                                                  Similarity
                                                                KLSMGSGLNLSEPN 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page; 190pp;
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2002US-0410260P.
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                                                                                                                                                              81.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                <u>.</u>
                                                                                                                                                              Score 60; DB 8;
Pred. No. 0.084;
                                                                                                                                   Mismatches
                                                                                                                                2
                                                                                                                                                                                         Length 609;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shen F;
SJ, Elder LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kirton ES;
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KLSMGSGLNLSEPN 15

Query Match Best Local S Matches 12

Similarity

Score 60; DB 8; Pred. No. 0.087; 0; Mismatches

Length 627

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0

Gaps

0

Conservative

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RESULT 11
ABM63781
ID ABM633
XX ABM63
XX ABM63
XX ABM63
XX ABM63
XX ABM63
XX PAM63
X
                                                        The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dith) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders, associated autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mooney EM,
Stevens KA,
Peralta CH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or in gene mapping.
   Sequence 627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 27; Page; 190pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lagace RE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-SEP-2003; 2003WO-US028227.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004-329368/30.
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S, Shi X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Delegeane AM, Panesar IS, Banville SC, Reddy TP; Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton mg M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gi Shi X, Suarez CJ;
       A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human
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Suchorolski MT, Altus CM, Pitts SJ, Elder LN
Suchorolski MT, Banville SC, Reddy TA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnostic and therapeutic polynucleotide; dithp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shen r;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kirton ES;
JL, Gietzen D;
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                                                                                Query Match
Best Local S
Matches 12
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Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES;
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gletzen
Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosting a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or in gene mapping.
                                                                                                                                                                     Sequence 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy; human diagnostic and therapeutic polynucleotide; dithp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human diagnostic and therapeutic pprotein SEQ ID NO:4028
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DB; ACN42431.
                                                                                  12; Conserv
                                        2 KLSMGSGLNLSEPN
KLCMGSGLNLCEPN 476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page; 190pp; English.
                                                                                     Conservative
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                                                                                                                                                                                                              www.wipo.int/pct/en/sequences/listing.htm
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                                                                                                    81.1%;
85.7%;
                                          15
                                                                                     <u>,</u>
                                                                                                       Score 60; DB 8;
Pred. No. 0.087;
                                                                                     Mismatches
                                                                                     2; Indels
                                                                                                                             Length 628;
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                                                                                Gaps
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RESULT 13

RESULT 14 ABM83778 ID ABM83 XX

ABM83778 standard; protein; 643

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ADTO8174

ADTO8174

AC ADTO
XX X ADTO
DT 30-[

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                                                                                                                                  Matches
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                 Sequence 635 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New human metal binding proteins (MBP), useful for diagnosing, treating and preventing diseases or conditions associated with the aberrant MBP expression e.g. cancer, AIDS, epilepsy, or infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jiang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MBP; metal binding protein; cytostatic; antiarteriosclerotic; anti-HIV; anticonvulsant; nootropic; neuroprotective; cerebroprotective; anti-HIV; antiallergic; antiinflammatory; thyromimetic; transgenic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADT08174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 3; 161pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-MAR-2003; 2003US-0457538P
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                                                                                                                                                                  Local
452
                                                                                                                                  12;
                                                            2 KLSMGSGLNLSEPN 15
                                                                                                                                                              Similarity
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                                                                                                                                  Conservative
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                                                                                                                                                              81.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hafalia AJA,
                                                                                                                                  0
                                                                                                                                  Score 60; DB 8;
Pred. No. 0.088;
0; Mismatches
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                                                                          RESULT 15
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                                                                                                                                                                                                                                                                                                                                      The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be cCC used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine CC disorder, neurological disorder, developmental disorders, or CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The sequence represents a dithp protein of the CC invention. Note: The sequence represents a dithp protein of the CC invention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention. Note: The sequence represents a dithp protein of the CC convention.
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Matches 12
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                    ABM83783;
                                                      ABM83783 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 27; Page; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
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12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                  Local Similarity
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Pred. No. 0.09;
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JL, Gietzen
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Mooney EM, Lanciu...
Stevens KA, Blanciu...
Peralta CH, Anderso
Peralta CH, Spiro P/
Padce RE, Spi
                                                                                                                                                     Matches
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Xu Y, Kwong M, Po
                                                                                                                                                                                                                                                                                                  Sequence 645 AA;
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DB; ACN42435.
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                                                                      2 KLSMGSGLNLSEPN 15
                                                                                                                                                                                    Similarity
KLCMGSGLNLCEPN
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                                                                                                                                                     Conservative
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Suchorolski MT, Altus CM, Pitts SJ, Elder L
Suchorolski MT, Banville SC, Reddy TP;
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Search completed: June 13, Job time : 97.0588 secs

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00016BOAD; 3.; Trepat. A for human	R; Bark, I; Schaefter, E; Sidoll, A; Baralle, F.E; Conen, G.N.; Zakin, M.M. Proc. Natl. Acad. Sci. U.S.A. 82, 3149-3153, 1985 A; Title: Organization of the human transferrin gene: direct evidence that it o A; Reference number: A94044; MUID:85216459; PMID:3858812 A; Accession: A94044 A; Molecule type: DNA A; Residues: 73-263, 'E', 265-328, 'N', 330-562 < PAR> A; Residues: 73-263, 'E', 265-328, 'N', 330-562 < PAR> A; Cross-references: UNIPARC:UPI0000174404; EMBL:M11361 R; Adrian, G.S.; Korinek, B.W.; Bowman, B.H.; Yang, F. Gene 49, 167-175, 1986 Gene 49, 167-175, 1986 A; Title: The human transferrin gene: 5' region contains conserved sequences wh A; Reference number: A29090; MUID:87192006; PMID:3106157 A; Accession: A29090	RESULT 1 TPHUD TPHUD transferrin precursor [validated] - human N;Alternate names: siderophilin C;Species: Homo sapiens (man) C;Apecies: Homo sapiens (man) C;Apecies: Homo sapiens (man) C;Apecies: Homo sapiens (man) C;Apecies: Lum, J.B.; McGill, J.R.; Moore, C.M.; Naylor, S.L.; van Bragt, P.H Proc. Natl. Acad. Sci. U.S.A. 81, 2752-2756, 1984 A;Title: Human transferrin: cDNA characterization and chromosomal localization A;Reference number: A20981; MUID:84194084; PMID:6585826 A;Contents: variant C A;Accession: A20981 A;Molecule type: mRNA A;Residues: 1-698 cyAN> A;Cross-references: UNIPROT:P02787; UNIPARC:UPI000002F089; EMBL:M12530; NID:93 A;Cross-references: UNIPROT:P02787; UNIPARC:UPI000002F089; EMBL:M12530; NID:93 A;Title: The authors translated the codon CAA for residue 203 as Glu R;MacGillivray, R.T.A.; Mendez, E.; Shewale, J.G.; Sinha, S.K.; Lineback-Zins, J. Biol. Chem. 258, 3543-3553, 1983 A;Title: The primary structure of human serum transferrin. The structures of s A;Reference number: A92417 A;Rolecule type: protein A;Residues: 20-263, 'E',265-328,'N',330-379,'SD',382-435,'D',437-557,'T',559-56 A;Cross-references: UNIPARC:UPI0000174483 A;Note: the sequence shown is the predominant electrophoretic genetic variant	ALIGNMENTS	34 38 51.4 1021 2 T05901 hypothetical prote 53 38 51.4 2212 2 A41098 calcium channel prote 36 37.5 50.7 431 2 F95236 PTS system, IIC co 37 37.5 50.7 431 2 H98100 hypothetical prote 53 37.5 50.7 431 2 H98100 hypothetical prote 63 37 50.0 106 2 T20860 hypothetical prote 64 37 50.0 124 2 B27632 T-cell receptor al 64 37 50.0 282 2 F71973 hypothetical prote 64 37 50.0 325 2 T35318 probable integrase 64 37 50.0 335 2 C95118 hypothetical prote 64 37 50.0 335 2 A97988 hypothetical prote 64 37 50.0 335 2 A97988 hypothetical prote 64 37 50.0 335 2 A97988 hypothetical prote 64 37 50.0 344 1 RWRTC2
EMBL:M15673 J.S.; Zakin, M.M.; Ka, transferrin. B; PIDN:AAA61142.1; PI)	e that it originated e that it originated equences which match	hange 09-Jul-2004 9; 16313; I54011; 168160; A03, 9; 163133; I54011; 168160; A03, S.L.; van Bragt, P.H.; Baldwin mosomal localization. EMBL:M12530; NID:g339452; PID) 03 as Glu 03 as Glu 03 S.K.; Lineback-Zins, J.; Brew, The structures of seven cyano; 1,437-557,'T',559-560,'P',562- tic genetic variant (C or TfC)		tical prote stem, IIC co stical prote receptor al rical prote receptor al rical prote le integrase le dehydroge stical prote stical prote

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Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli
A;Reference number: A64720; MUID:97426617; PMID:9278503
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A;Title: Changes in brain gene expressions shared by scrapie and Alzheimer disease. A;Reference number: I48174; MUID:89386721; PMID:2780570
A;Accession: I63133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A,Description: binds iron for delivery into cells (;Superfamily: transferrin; transferrin repeat homology C;Keywords: duplication; glycoprotein; iron transport; metal binding; plasma (;Keywords: duplication; glycoprotein; iron transport; metal binding; plasma (;Leywords: dipal sequence #status predicted <SIG> F;20-698/Product: transferrin #status experimental <MAT> F;20-350/Domain: transferrin repeat homology <TRH1> F;356-686/Domain: transferrin repeat homology <TRH1> F;356-686/Domain: transferrin repeat homology <TRH2> F;38-67,38-58,137-213,156-350,177-193,180-196,190-198,246-260,358-615,364-396,374-387,42 F;432,630/Binding site: carbohydrate (Asn) (covalent) #status experimental
                                                                                                              hypothetical protein b2850 - Escherichia coli (strain K-12) C;Species: Escherichia coli C;Species: Escherichia coli C;Species: Escherichia coli C;Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 09-Jul-2004 C;Accession: C65068 R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; J. Rose, D.J.; Mau, B.; Shao, Y.
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A;Title: Complete structure of the human transferrin gene. Comparison with analogous chi A;Reference number: I54011; MUID:88056305; PMID:3678832
A;Accession: I54011
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-72 <RE3>
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A; Introns: 15/1;
C; Function:
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A;Cross-references: GDB:120432; OMIM:190000
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A;Status: translated from GB/EMBL/DDBJ
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 564-698 <RE4>
A;Residues: 564-698 <RE4>
A;Cross-references: UNIPARC:UPI000016B0AE; GB:M17614; NID:g339483; PIDN:AAA61148.1; PID:C;Comment: Each of the two repetitive domains binds a ferric ion and a bicarbonate anion C;Cennetics:
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A; Residues: 636-696 < RE2>
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A,T1tle: The complete amino acid sequence of human serum transferrin.
A,Reference number: A93911, MUID:82222166; PMID:6953407
A,Contents: annotation; disulfide bonds
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A;Title: A cloned gene for human transferrin.
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C; Genetics:
A; Gene: ECs3707
C; Superfamily: 1
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                                                                                                                                                                                                                                               A;Title: Genome sequence of A;Reference number: A85480; A;Accession: G85937
                                                                                                                                                                                                                                                                                                           R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glassiller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein Z4171 [imported] - Escherichia coli (strain O157:H7, substrain C;Species: Escherichia coli (c;Species: Escherichia coli (c;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004 C;Accession: G85937
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A;Molecule type: DNA
A;Residues: 1-148 <BLAT>
                                                                                   A; Gene: Z4171
C; Superfamily:
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A; Residues: 1-163 < HAY>
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KLSMGSGLNLSE 13

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RESULT 7
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N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
A;Variety: isolate 09D09HCC
C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 09-Jul-2004
C;Accession: T13469
R;Takahashi, K:, Akahane, Y.; Hino, K.; Ohta, Y.; Mishiro, S.
Arch. Virol. 143, 2313-2326, 1998
A;Title: Hepatitis B virus genomic sequence in the circulation of hepatocell
A;Reference number: Z17684; MUID:99129050; PMID:9930189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N;Contains: major surface antigen; middle surface antigen C;Species: hepatitis B virus, HBV A;Variety: isolate 29Y11HCC C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 31-Dec-2004 C;Accession: T13474 R;Takahashi, X; Akahane, Y; Hino, K.; Ohta, Y.; Mishiro, S. Arch. Virol. 143, 2313-2326, 1998
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N;Alternate names: envelope protein; HBs antigen
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Keywords: glycoprotein; surface antigen
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Pred. No.
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Pred. No. 21;
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N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A03705; 504569; JQ2107; PQ0608
R;Ono, Y:; Onda, H:; Sasada, R:; Igarashi, K:; Sugino, Y:; Nishioka, K:
Nucleic Acids Res. 11, 1747-1757, 1983
A;Title: The complete nucleotide sequences of the cloned hepatitis B virus DNA; subtype
A;Reference number: A93460; MUID:83168919; PMID:6300776
A;Accession: A03705
                        R;Mukaide, M.; Kumazawa, T.; Hoshi, A.; Kawaguchi, R.; Hikiji, K. Nucleic Acids Res. 20, 6105, 1992
A;Title: The complete nucleotide sequence of hepatitis B virus, starterence number: S35527; MUID:93096607; PMID:1461746
A;Accession: S35528
                                                                                                                                                                                                                                                                                             surface antigen - hepatitis B virus (subtype adr)
N;Alternate names: envelope protein; HBs antigen
N;Contains: surface antigen pre-S1 (large envelope protein); surface antigen
C;Species: hepatitis B virus, HBV
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A; Molecule type: DNA
A; Residues: 275-354 < NO2>
A; Cross-references: UNIPARC: UPI000017496D
A; Cross-references: subtype adrq+, Bau
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Keywords: glycoprotein; surface antigen (gene pre-S2/S) #status predicted F;120-400/Product: major surface antigen (gene S) #status predicted cNSA>
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A; Residues: 1-50,'r,52-66,'YP',69-129,'V',131-142,'P',144-164,'S',166-176,'S',178-338,'
A; Cross-references: UNIPARC: UPIO00017496C; EMBL: X14193
A; Cross-references: UNIPARC: UPIO00017496C; EMBL: X14193
A; Cross-references: UNIPARC: UPIO00017496C; EMBL: X14193
Cen. Virol. 74, 1341-1348, 1993
J. Gen. Virol. 74, 1341-1348, 1993
J. Gen. Virol. 74, 1341-1348, 1993
                                                                                                                                                                                                 A;Variety: subtype adr
C;Date: 09-Dec-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S35528
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A;Experimental source: genogroup C, subtype adr, strain pBRHBadr4
R;Norder, H.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 3141-3145, 1992
A;Title: Molecular basis of hepatitis B virus serotype variations within the four major A;Reference number: PQ0453; MUID:93107848; PMID:1469353
A;Accession: PQ0608
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A; Residues: 175-400 < NOR>
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A; Residues: 1-400 < ONO>
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N;Contains: major surface antigen; middle surface antig
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Status: nucleic acid sequence not shown; translation not
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A;Variety: subtype adr
C;Date: 07-Sep-1994 #sequence_revision 10-Nov-1995 #text_change
C;Accession: $43492; PN0601; PN0602
C;Accession: $43492; PN0601; PN0602
R;Loncarevic, I.F.; Zentgraf, H.; Schroeder, C.H.
Nucleic Acids Res. 18, 4940, 1990
A;Title: Sequence of a replication competent hepatitis B virus
A;Reference number: $12598; MUID:90370503; PMID:2395664
A;Accession: $43492
A;Accession: $43492
A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-445 <LON>
A;Residues: 1-445 <LONA
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Alintrons: 165/2
C:Superfamily: hepatitis B virus surface antigen
C:Keywords: Surface antigen
F:1-400/Product: surface antigen pre-S1 (large envelope protein)
F:1-400/Product: surface antigen pre-S1 (large envelope protein)
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                                                                                                                                                                                                                                                                                                      F;1-164/Domain: pre-S1 domain #status predicted <PRE1>
F;165-445/Product: surface antigen pre-S2 (middle envelope protein)
F;165-219/Domain: pre-S2 domain #status predicted <PRE2>
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A;Residues: 165-196 <MI2>
A;Cross-references: UNIPARC:UPI0000178455
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A;Gene: S
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;Accession: PN0602
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                                                                                                                                               Similarity
9; Conserv
                       SKPRQGMGTNLSVPN
                                                                                SKLSMGSGLNLSEPN 15
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                                                                                                                                               54.1%;
llarity 60.0%;
Conservative
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                       65
                                                                                                                                               Score 40; DB Pred. No. 25; 0; Mismatches
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A; Description: The sequence A: Reference number: Z20429
                                                       C;Accession: T27845
R;Favello, T.
submitted to the EMBL Data Library, November 1995
                                                                                                                        hypothetical protein ZK402.5 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;pate: 15-Oct-1999 #t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: DNA
A;Restdues: 1-448 <TOM>
A;Crestdues: 1-448 <TOM>
A;Cross-references: UNIPROT:P56005; UNIPARC:UPI0000135EDF; GB:AE000621; GB:AE000511;
C;Superfamily: signal recognition particle 54K protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nature 388, 539-547, 1997
A; Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A; Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A; Reference number: A64520; MUID:97394467; PMID:9252185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R.D., Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKenney son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey, L.
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H64663
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A;Gene: ffh
C;Superfamily: signal recognition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable signal C; Species: Helio
A; Reference number: A; Accession: T27845
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      signal recognition particle protein - Helicobacter pylori (strain 26695)
C;Species: Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q9ZK62; UNIPARC:UPI0000135EDE;
A;Experimental source: strain J99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: Genomic sequence comparison of two unrelated isolates of the human gastric pathous, Reference number: A71800; MUID:99120557; PMID:9923682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Species: Helicobact A; Variety: strain J99
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Best Local
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7; Conserve
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5; Mismatc
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N;Contains: Bullion Fig. HBV
C;Species: hepatitis B virus, HBV
A;Variety: subtype adw2
C;Date: 31-Dec-1993 #sequence_revision 22-Oct-1999 #text_change 09-Jul-2004
C;Accession: S47411; JQ1575
C;Accession: S47411; JQ1575
R;Plucienniczak, A.
submitted to the EMBL Data Library, August 1994
submitted to the EMBL Data Cloning and sequencing of two complete genomes of
                  A,Description: Molecular cloning and sequencing of two complete genomes of polish isolat A;Reference number: $47404
A;Reference number: $47401
A;Accession: $47411
A;Molecule type: DNA
A;Residues: 1-400 <PLU>
A;Cross-references: UNIFROT:067896; UNIPARC:UPI00000F5BAF; EMBL:235717; NID:g527440; PII
A;Cross-references: UNIFROT:067896; UNIPARC:UPI00000F5BAF; EMBL:235717; NID:g527440; PII
A;Experimental source: subtype adw2
R;Norder, H.; Hammas, B.; Loefdahl, S.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 1201-1208, 1992
A;Title: Comparison of the amino acid sequences of nine different serotypes of hepatitis A;Reference number: JQ1570; MUID:92268879; PMID:1588323
A;Accession: JQ1575
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A;Experimental source: strain N315
C;Genetics:
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C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004
C;Accession: C89991
R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.;
ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu,
C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.
Lancet 357, 1225-1240, 2001
A;Title: Whole genome sequencing of meticillin-resistant Stapylococcus aureus.
A;Reference number: A89758; MUID:21311952; PMID:11418146
A;Accession: C89991
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A;Molecule type: DNA
A;Residues: 1-516 <FAV>
A;Cross-references: UNIPROT:Q23477; UNIPARC:UPI00000783D4; EMBL:U40956; PIDN:AAA81756.1
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        N;Alternate names: envelope protein; HBs antigen N;Contains: surface antigen pre-S1 (large envelope C;Species: hepatitis B virus, HBV
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C; Superfamily:
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A;Introns: 28/2; 122/2; 160/2; 248/2; 420/3
C;Superfamily: Caenorhabditis elegans hypothetical
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A; Residues: 1-338 < KUR>
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type: DNA
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Pred. No.
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C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
C;Keywords: glycoprotein; surface antigen
F;120-400/Product: middle surface antigen (gene pre-2/S) #status predicted
F;175-400/Product: major surface antigen (gene S) #status predicted <MSA>
F;15,123,177/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        large surface antigen - hepatitis B virus (subtype adw, strain 991)
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Date: 30-Sep_1991 #sequence_revision 30-Sep-1991 #text_change 09-Jul-2004
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A;Cross-references: UNIPARC:UPI00000002E4; GB:X75666; NID:g416074; PIDN:CAA53362.1; A;Experimental source: subtype adw2, strain P6
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F;1-119/Domain: pre-S1 domain #status predicted <PRE1>
F;120-400/broduct: surface antigen pre-S2 (middle envelope protein) #status predicted F;120-174/Domain: pre-S2 domain #status predicted <PRE2>
F;175-400/broduct: surface antigen S (small envelope protein) #status predicted <PSD>
F;177/Binding site: carbohydrate (Asn) (covalent) #status predicted
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C; Superfamily: h
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A; Residues: 1-400 < KOE>
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Pred. No.
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Gapop 10.0 , Gapext 0.5
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74
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1: uniprot_sprot:*
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           SKLSMGSGLNLSEPN 15
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   BB
TRFE HUMAN
Q53HZ6 HUMAN
Q53HZ6 HUMAN
Q58HZ7 PONPY
Q98952 HBV
Q943K CORYSA
Q54102 CANAL
Q45RA, STRFR
Q3Y1Z5 HAEFR
Q3Y1Z5 HAEFR
Q3Y1Z5 HAEFR
Q3Y1Z5 HAEFR
Q5SBE11 EMENI
Q5CSS2 CRYPV
Q4SQN4 TETNG
Q5SNH2 CANAL
PDK1L BRARE
Q5NH2 CANAL
PDK1L BRARE
Q5NH2 TENNAN
Q5TQJZ HUMAN
Q5TQJZ ANGGA
Q43GK1 GCANGA
Q41GK1 CANGA
Q2UC93 ANGGA
Q41GK1 CANGA
Q2UC93 ANGGA
Q41GK1 CANGA
Q2UC93 ANGGA
Q41GK1 CANGA
Q2UC93 TAFFU
Q3YY03 SHISS
YGEF ECOLI
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Q9D013 MOUSE
Q8X6HB ECO57
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P02787 homo sapien Q53h26 homo sapien Q57917 pongo pygma Q9e952 hepatitis b Q943k6 oryza sativ Q5alq2 candida alb Q8j212 candida alb Q8j212 candida alb Q8j212 candida alb Q5css5 cryptospori Q4sqn4 tetraodon n Q59h2 brachydanio Q8nc87 homo sapien Q9uif6 homo sapien Q5tq12 anopheles g Q43gk1 chlorobium Q6fmk0 candida gla Q2uc93 aspergillus Q8cs9 kluyveromyc Q4r5m1 macaca fasc Q8u379 pyrococcus Q8j2997 cryptospori Q3yy09 shigella so Q46786 escherichia Q3w4678 mescherichia Q9w497 hepatitis b
                                                                                                                                                                                                                                                                                Description
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ALIGNMENTS

NUCLEOTIDE SEQUENCE MEDLINE=20563920; p Beutler E., Gelbart Fairbanks V.F.; molecular characte Blood 96:4071-4074([5] NUCLEOTIDE SEQUENCE Rieder M.J., Carrin Ahearn M.O., Kuldan Nickerson D.A.; "SeartleSNPs WHUBI FHCRC, Seattle, WA Submitted (MAY-2003	NUCLEOTIDE SEQUENCE. MEDLINE=88056305; PubMed=3678832. Schaeffer E., Lucero M.A., Jelts. Chambon P., Cohen G.N., Zakin M.I. "Complete structure of the human analogous chicken gene and human Gene 56:109-116(1987). [3] NUCLEOTIDE SEQUENCE. TISSUE=Liver; MEDLINE=92231399; PubMed=1809186, MEDLINE=9231399; PubMed=1809186, Hershberger C.L., Larson J.L., Alwilliams P., Dehoff B., Dunn P., "A cloned gene for human transferent of the second of the s	HOMO. (1) (1) NCBI_TAXID=9606; (1) NUCLEOTIDE SEQUENCE, AND VARI MEDLINE=84194084; PubMed=6585 Yang F., Lum J.B., McGill J.R. Yan Bragt P.H., Baldwin W.D., "Human transferrin: cDNA char "localization."; Proc. Natl. Acad. Sci. U.S.A.	RESULT 1 TRFE HUMAN ID TOT-87; 043890; Q9NQB8; Q9UHVO; DT 21-JUL-1986, integrated into UniProtKB/Swiss- DT 01-0CT-1989, sequence version 2. DT 07-MAR-2006, entry version 84. DT 07-MAR-2006, entry version 84. DE Serotransferrin precursor (Transferrin) (Side DE binding globulin). GN Name=TF; ORFMames=FR01400; OS Homo sapiens (Human). OC Eukaryota; Metazoa; Chordata; Craniata; Verte OC Mammalia; Butheria; Euarchontoglires; Primate
, AND VARIANT ATRANSFERRINEMIA PRO-477. ubMed=11110675; T., Lee P.L., Trevino R., Fernandez M.A., rization of a case of atransferrinemia."; 2000). [GENOMIC DNA]. [GENOMIC DNA]. gton D.P., da Ponte S.H., Hastings N.C., ek S.A., Rajkumar N., Toth E.J., Yi Q., HI.66682 program for genomic applications, UW- (URL: http://pga.gs.washington.edu)."; to the EMBL/GenBank/DDBJ databases.	DOI=10.1016/0378-1119(87)90163-6; th JM., Py MC., Levin M.J., transferrin gene. Comparison with pseudogene."; cnold B., Rosteck P.R. Jr., o'Neal K.L., Riemen M.W., Tice P.A.; trin."; (1991).	2; TF*CHI AND TF*D1. C.M., Naylor S.L., .H.; ion and chromosomal 2756(1984).	698 AA. KB/Swiss-Prot. in) (Siderophilin) (Beta-1-metal- tta; Vertebrata; Euteleostomi; primates; Catarrhini; Hominidae;

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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Warusina K., Foshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Hitting M., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schemtz J., Myers R.M.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalbka U., Smailus D.E.,
Tend mouse cDNA secmences "."
"Generation and initial analysis of more than 15,000 full-length human
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NUCLEOTIDE SE
TISSUE=Brain;
         Gene
                          Adrian G.S., Korinek B.W., Bowman B.H., Yang F
"The human transferrin gene: 5' region contain
which match the control elements regulated by
glucocorticoids and acute phase reaction.";
Gene 49:167-175(1986).
                                                                                                                                                                                                                                                                                                                          [10]
NUCLEOTIDE SEQUENCE OF 73-698.
MEDLINE=85216459; PubMed=3888812;
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                                                                                                                                                                                   WUCLEOTIDE SEQUENCE OF 1.14.
MEDLINE=87066744; PubMed=3786138;
Lucero M.A., Schaeffer E., Cohen G.N., 2
"The 5' region of the human transferrin
                                                                                                                                                                                                                                                                                                                                                                                                             seven cyanogen
structure.";
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Lineback-Zins J., Brew K.;
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Zakin M.M., Kahn A.;
"Molecular cloning and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Functional prediction of the coding sequences of 33 new by analysis of cDNA clones from human fetal liver."; Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                     originated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEIN SEQUENCE OF 20-698.
MEDLINE=83160878; PubMed=6833213;
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                                                                                                                                                                                                                                                                                    "Organization of the human transferrin gene: originated by gene duplication.";
                                                                                                                                                                                                                                                                                                                   Zakin M.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biochem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transferrin."
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1. Acad. Sci. U.S.A. 99:16899-16903(2002)
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hn A.;
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PubMed=3106157; DOI=10.1016/0378-1119(86)90277-5;

nek B.W., Bowman B.H., Yang F.;

errin gene: 5' region contains conserved sequences
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X-RAY CRYSTALLOGRAPHY (1.6 ANGSTROMS) OF 22-350.
MEDLLNE=98272665; PubMed=9609685; DOI=10.1021/bi980355j;
Macgillivray R.T.A., Moore S.A., Chen J., Aherson B.F.,
Luo Y., Bewley M.C., Smith C.A., Murphy M.E.P., Wang Y.,
Woodworth R.C., Brayer G.D., Baker E.N.;
"Two high-resolution crystal structures of the recombinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Namekata K., Oyama F., Imagawa M., "Human transferrin (Tf): a single Cl or Tf C2 variant.";
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DOI=10.1002/1097.4547(20000815)61:4x388::AID-JNR5>3.0.CO;2-Q;
DOI=10.1002/1097.4547(20000815)61:4x388::AID-JNR5>3.0.CO;2-Q;
de Arriba Zerpa G.A., Saleh M.-C., Fernandez P.M., Guillou F.,
Espinosa de los Monteros A., de Vellis J., Zakin M.M., Baron B
"Alternative splicing prevents transferrin secretion during
differentiation of a human oligodendrocyte cell line.";
X-RAY CRYSTALLOGRAPHY (2.2 ANGSTROMS) OF 22-350.
MEDLINE=9843436; PubMed=9760232; DOI=10.1021/bi9812064;
Jeffrey P.D., Bewley M.C., Macgillivray R.T.A., Mason A.
Woodworth R.C., Baker E.N.;
                                                                                                                                                                                                                                     "Human plasma N-glycoproteome analysis by immunoaffinity hydrazide chemistry, and mass spectrometry.";
J. Proteome Res. 4:2070-2080(2005).
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                                                                                                                                                                                                                                                                                                                                                                               Woodworth R.C., Mason A.B., Funk W.D., McGillivray R.T.A.; "Expression and initial characterization of five site-directed mutants of the N-terminal half-molecule of human transferrin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=82222166; PubMed=6953407; McGillivray R.T.A., Mendez E., Sinha Lineback-Zins J., Brow K.;
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Ershova E.S., Egorov T.A., Mus
"The major protein expression
database of human heart.";
                                                                                                                                                                                                                                                                                                                                                               Biochemistry 30:10824-10829(1991)
                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=92031536; PubMed=1932003;
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Duguid J.R., Bohmont C.W., Liu N.G., Tourtellotte W.
"Changes in brain gene expression shared by scrapie
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RESULT 2
Q53H26 HUMAN
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Q53H26;
DT 24-MAY-2005, sequence version 1.
DT 24-MAY-2005, sequence version 1.
DT 24-MAY-2005, sequence version 1.
DT 07-FEB-2006, entry version 4.
DE Transferrin variant (Fragment).
OS Homo sapiens (Human)
OC Mammalia; Eutheria; Euarchontoglire OC Homo.
OX NCBI_TaxID=9606;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RC TISSUE=Liver;
RX MEDLINE=94171032; PubMed=8125298; D
RA Maruyama K., Sugano S.;
RN [2]
RN NUCLEOTIDE SEQUENCE.
RC TISSUE=Liver;
RX MEDLINE=9803896; PubMed=9373149; D
RX MEDLINE=9803896; PubMed=9373149; D
RX SUZUKİ Y., Yoshitomo K., Maruyama K
RC TISSUE=Liver;
RX MEDLINE=9803896; PubMed=9373149; D
RX MEDLINE=9803896; PubMed=9373149; D
RX MCLEOTIDE SEQUENCE.
RC TISSUE=Liver;
RX SUZUKİ Y., Yoshitomo K., Maruyama K
RT "Construction and characterization end-enriched cDNA library.";
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RC TISSUE=Liver;
RC TISSUE=Liver;
RC Gene 200:149-156(1997).
RN [3]
RP NUCLEOTIDE SEQUENCE.
RC TISSUE=Liver;
RC TISSUE=Liver;
RC GOUNGENCE.
TOOKSANA A., Yokoyama S.;
RC SUZUKİ Y., Sugano S., Totoki Y., To RA Tanaka A., Yokoyama S.;
RC SUZUKİ Y., Sugano S., Totoki Y., To RA Tanaka A., Yokoyama S.;
RC SUZUKİ Y., Sugano S., Totoki Y., To RA Tanaka A., Yokoyama S.;
RC GOUNGESTE; BAD96475.1; - mRNA SMR; Q53H26; 23-346.
DR EMBL; AKZ22755; BAD96475.1; - mRNA SMR; Q53H26; 23-346.
DR EMBL; AKZ22755; BAD96475.1; - mRNA SMR; Q53H26; 23-346.
DR Ensembl; ENSGOONO091513; Homo sapi GO; GO:0006879; P:icon ion transpor InterPro; JPRO01156; Peptidase S60; PIRMINES PERVIN.
DR PANTHER; PS00206; TRANSFERRIN.
DR PANTHER; PS00206; TRANSFERRIN.
2 PROSITE; PS00206; TRANSFERRIN.
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SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
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PROSITE; PS00207; TRANSFERRIN 3; 2.
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Tanaka A., Yokoyama S.;
Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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GO:0005576; C:extracellular region; IEA.
GO:0008199; F:ferric iron binding; IEA.
GO:0006879; P:iron ion homeostasts; IEA.
GO:0006876; P:iron ion transport; IEA.
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Pred. No. 0.09;
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Pred. No. 0.09;
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DT 07-FEB-2006.
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Best Local
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EMBL; CR859370; CAH91543.1; -; mRNA.
SMR; Q5R9L7; 23-346.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
Interpro; IPRO01156; Peptidase_S60;
PANTHER; PTHR11485; Peptidase_S60; 1.
Pfam; PF00405; Transferrin; 2.
                                                                                                                                                                                                                01-MAR-2001, integrated into UniProtKB/TrEMBL 01-MAR-2001, sequence version 1. 07-FEB-2006, entry version 16. Large surface antigen. Hepatitis B virus (HBV).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 2; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
NUCLEOTIDE SEQUENCE.
MEDLINE=20528423; PubMed=11074474;

MEDLINE=20528423; PubMed=11074474;

DOI=10.1002/1096-9071(200012)62:44456::AID-JMV10>3.0.CO;2-3;

Sugauchi F., Mizokami M., Orito E., Ohno T., Kato H., Maki M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bloecker H., Boecher M., Brandt P., Mewes H.W., Weil B., Osanger A., Fobo G., Han M., Wiemann S.; Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-DEC-2004, integrated into UniPro
21-DEC-2004, sequence version 1.
07-FEB-2006, entry version 6.
Hypothetical protein DKFZp459H0229
                                                                                                                                                                                          Viruses; Retro-transcribing viruses; Hepadnaviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pongo pygmaeus (Orangutan).
Eukaryota; Metazoa; Chordat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The German cDNA Consortium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KLCMGSGPHLCEPN 527
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Putative SUT1 protein (Sodium/lithium tolerance protein).
Name=P0031D11.42; Synonyms=P0408C03.17;
Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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Best Local S
Matches 10
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Jones T., Federspiel N.A., Chibana H., Dungan J., K.
Magee B.B., Newport G., Thorstenson Y.R., Agabian N
Davis R.W., Scherer S.;
"The diploid genome sequence of Candida albicans.";
proc. Natl. Acad. Sci. U.S.A. 101:7329-7334 (2004).
-!- CAUTION: The sequence shown here is derived from
EMBL/GenBank/DDBJ whole genome shotgun (WGS) en
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26-APR-2005, sequence version 1.
07-FEB-2006, entry version 7.
Hypothetical protein GPR1.
Name-GPR1, ORFNames=CaO19.1944, CaO19.9499;
                                                                                                                                                                                                                          EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Candida albicans SC5314.
Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
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EMBL; AP003241; BAB93215.1; -; Genomic_DNA.
EMBL; AY672599; AAT81168.1; -; Genomic_DNA.
Gramene; Q943K6; -.
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"Posttranslational modification of the rice SLT proteins for chaperone activity under stress conditions.";
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
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                                                                                                               PROSITE; PS00018; EF
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q5A1Q2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=237561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
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                                                                                                                                          AACQ01000082; EAK96633.1; -; Genomic_DNA.
AACQ01000081; EAK96692.1; -; Genomic_DNA.
Pro; IPR002048; EF hand Ca bd.
TE; PS00018; EF_HAND_1; UNKNOWN_1.
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ene family: a new subclass
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                                                                           92369 MW;
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71.4%;
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Pred. No.
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                                                                               ACESIC1C37711823 CRC64;
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H., Hahn J.H., Kim H.-
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Length 823;
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H.-I.,
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13-SEP-2005, e
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Hypothetical
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Q8J2T2;
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SEQUENCE
                                                                                                                                                             Baltz R.H.;
"The lipopeptide antibiotic
Streptomyces fradiae.";
Streptomyces fradiae.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein.
Streptomyces fradiae.
Bacteria, Actinobacteria;
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GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR002048; EF hand Ca bd.
PROSITE; PS00018; EF_HAND_1; UNKNOWN_1.
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharotomycetales; mitosporic Saccharomycetales; Candida.

NCBI_TaxID=5476;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-2003, integrated into U
01-MAR-2003, sequence version
07-FEB-2006, entry version 12.
                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
STRAIN=NRRL18158;
                                                                                                                                                                                                                                                                                                                                                                                                                            Streptomycineae;
NCBI_TaxID=1906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and hypha formation in the pathogenic Eukaryot. Cell 3:919-931(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
PubMed=15302825; DOI=10
Miwa T., Takagi Y., Shi
Kumagai H., Tamaki H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G-protein coupled receptor.
                                                                                                                                                                                                                                                                                     Miao V., Brost R., Chapple J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Gprl,
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                                                                                                                                                  J. Ind. Microbiol. Biotechnol. 0:0-0(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CANAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AB084519;
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SKLSMGSGLNLSEP 14
| :|||:|:| :||
SPMSMGAGINWTEP 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPMSMGAGINWTEP 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    823 AA; 92341 MW;
                                           by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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Y., Shinozaki M., Yun C.W., Schell W.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Streptomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62.2%;
57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Actinobacteridae; Actinomycetales; ycetaceae; Streptomyces.
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Pred. No. 43;
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                                                                                                                                                                                                                                                                                     K., Coeffet-Le Gal M.-F.,
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ic fungus Candida albicans."
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Best Local
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Pfam; PF00551; Abhydrolase_1; 1.
26-APR-2005, integrate
26-APR-2005, sequence
07-MAR-2006, entry ver
Hypothetical protein.
ORFNames=AN1219.2;
                                                                                                                                                                                                                                             EMBL; DQ127937; AAZ67641.1; -;
NON TER 1 1
SEQUENCE 444 AA; 49811 MW;
                                                                                                                                                                                                                                                                                                                                     Melnikow E., Dornan S., Sargent C., Duszenko M., Evans G., Guser P.M., Ullrich H.J.; Selzer P.M., Ullrich H.J.; "Microarray analysis of Haemophilus parasuis gene expression vitro growth conditions minicking the in vivo environment."; Submitted (JUL-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-SEP-2005, integrated into UniProt 27-SEP-2005, sequence version 1. 07-FEB-2006, entry version 3. Putative heme-hemopexin utilization
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PRINTS; PRO0412; EPOXHYDRLASE.
PROSITE; PS00560; CARBOXYPEPT SER HIS; 1.
Antibiotic biosynthesis; Hypothetical pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Haemophilus parasuis.
Bacteria, Proteobacteria,
Pasteurellaceae, Haemophi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                 Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=29775;
                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=738;
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                         entry version
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                                                   integrated into UniProtKB/TrEMBL
                                                                           PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    F:hydrolase activity; TEA.
F:serine carboxypeptidase activity; IEA.
P:antibiotic biosynthesis; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Haemophilus.
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                                                                                                                                                                                                           Score 45;
Pred. No.
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                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hetical protein.
598D45B0255F2C35 CRC64;
                                                                                                                                                                                                                                                                       Genomic_DNA
                                                                                                                                                                                                                                                 BB1A89DB46EF8565 CRC64;
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                             NUCLEOTIDE TYPE II;
STRAIN=IOWA type II;
PubMed=15044751; DOI=10.1126/science.1094786;
PubMed=15044751; DOI=10.1126/science.1094786;
Abrahamsen M.S., Templeton T.J., Enomoto S., Abrahante (Lancto C.A., Deng M., Liu C., Widmer G., Tzipori S., Bulancto C.A., Dear P.H., Konfortov B.A., Spriggs H.F., Bankier A.T., Dear P.H., Konfortov B.A., Spriggs H.F., Anantharaman V., Aravind L., Kapur V.;
Anantharaman V., Aravind L., Kapur V.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Sequencing or """;
fumigatus and A. oryzae.";
Nature 438:1105-1115(2005).
-I- CAUTION: The sequence shown here is derived in the sequence shown here is derived in the sequence should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be should be
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; Emericella.
NCBI_TaxID=227321;
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                                                                                                                                                                                                                                                                 parvum.";
Science 304:441-445(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cryptosporidium parvum.
Eukaryota; Alveolata; Apicomplexa;
Cryptosporidiidae; Cryptosporidium.
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                                                         by the UniProt Consortium, under the Creative Commons
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ckmen M., Spevak C.C., Clutterbuck J.,
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Attribution-NoDerivs License
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A Jaillon O., Aury J.-M., Brunet F., Petit J.-L., Stange-Thomann N.,

A Maucell E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,

A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,

Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,

A Dasilva C., Jubin C., Castelli V., Katinka M., Vacherie B.,

A Anthouard V., Jubin C., Cattolico L., Poulain J., De Berardinis V.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,

A Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,

A Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Bosak S.,

Kellis M., Volff J.-N., Guigo R., Zody M.C., Mesirov J.,

A Kellis M., Volff J.-N., Guigo R., Zody M.C., Mesirov J.,

A Kellis M., Volff J.-N., Guigo R., Zody M.C., Mesirov J.,

A Laudet V., Schachter V., Outerier F., Saurin W., Scarpelli C.,

Wincker P., Lander E.S., Weissenbach J., Roest Crollius H.;

"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
                                                                                                    Query Match
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
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19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 3.
Chromosome 17 SCAF14532, whole genome shotgun ORFNames=GSTENG00014297001;
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                                                                                                                                                                                                               SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the early vertebrate proto-karyotype."; Nature 431:946-957(2004).
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99528 MW; E
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Pred. No.
                                                                                                                                   Score 44;
Pred. No.
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Attribution-NoDerivs License
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QSSNH2;
QSSNH2;
26-APR-2005, integrated into UniPrc
26-APR-2005, sequence version 1.
07-FEB-2006, entry version 7.
Hypothetical protein.
ORFNames=Caol9.11674, Caol9.4198;
                                  EMBL;
                                                                                                                                                                                                                 NIH - Zebrafish Gene Collection (ZGC) project;
Submitted (OCT-2005) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: ATP + a protein = ADP + a phosphoprotein.
-!- SUBCELLULAR LOCATION: Nucleus (By similarity).
-!- SIMILARITY: Belongs to the Ser/Thr protein kinase family.
-!- CAUTION: Glu-37 is present instead of the conserved Lys which expected to be an active site residue. Lys-38 may fulfill that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Verrebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes; Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
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Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Serine/threonine-protein kinase pdik11 (EC 2.7.1.37). Name=pdik11; ORFNames=zgc:123209;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JAN-2006, integrated into UniProtKB/Swiss-Prot. 06-DEC-2005, sequence version 1. 07-MAR-2006, entry version 4.
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Q32PP3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The diploid genome sequence of Candida albicans.";
Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334(2004).
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=15123810; DOI=10.1073/pnas.0401648101; Jones T., Federspiel N.A., Chibana H., Dungan J., Magee B.B., Newport G., Thorstenson Y.R., Agabian Davis R.W., Scherer S.;
                                                                                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Ovary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE (LARGE
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     PROSITE; PS00107;
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                               BC108040; AAI08041.1; -; mRNA
ZDB-GENE-051113-236; zgc:12320
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PROTEIN_KINASE_ATP; FALSE_NEG
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Pred. No. 13;
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HSSP; Q9UIGO; 1F62.
Ensembl; ENSG00000123636; Homo sapiens.
GO; GO:0005515; F:protein binding; IEA.
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0008355; P:regulation of transcri
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01-OCT-2002, sequence version 1.
21-FEB-2006, entry version 17.
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PROSITE; PS00108; PROTEIN_KINASE_ST; 1.
ATP-binding; Kinase; Nuclear protein; Nucleotide-binding; Serine/threonine-protein kinase; Transferase.
CHAIN 1 341 Serine/threonine-protein kin
                                                                                                                                                                                                                                   SMART; SM00249; PHD; 1.
PROSITE; PS50016; ZF PHD_2; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria;
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                                                   SMGSGLNLSEPN
  SLGSGLGLSEGN
                                                                                                                                                                                               796 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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22
164
                                                                                                                                                                                               88463 MW;
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75.0%;
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ATP (By similarity).

ATP acceptor (By
                                                                                                                        Score 43; I
Pred. No. 1.
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Pred. No. 55;
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Search completed: June 13, 2006, 04:17:49
Job time: 124.103 secs

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43, Appl 45, Appl 46, Appl 47, Appl 48, Appl 48, Appl 48, Appl 48, Appl 48, Appl 49, Appl 41, Appl 42, Appl 43, Appl 43, Appl 44, Appl 45, Appl 47, Appl 47, Appl 47, Appl 47, Appl 48, Appl 49, Appl 49, Appl 41, Appl 41, Appl 42, Appl 43, Appl 44, Appl 45, Appl 46, Appl 47, Appl 48, Appl 49, Appl 49, Appl 49, Appl 40, Appl 40, Appl 41, Appl 42, Appl 43, Appl 44, Appl 45, Appl 46, Appl 47, Appl 48, Appl 49, Appl 49, Appl 40, Appl 41, Appl 41, Appl 42, Appl 43, Appl 44, Appl 45, Appl 46, Appl 47, Appl 48, Appl 49, Appl 49, Appl 40, Appl 40, Appl 41, Appl 42, Appl 43, Appl 44, Appl 45, Appl 46, Appl 47, Appl 48, Appl 49, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 41, Appl 42, Appl 43, Appl 45, Appl 46, Appl 47, Appl 48, Appl 49, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40, Appl 40,

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                         hits satisfying chosen parameters:
                                                                                                                                                                                             650591 seqs, 87530628 residues
                                                                                                                                                                                                                                                                                                                                                               GenCore version 5.1.9 Copyright (c) 1993 - 2006 Biocceleration Ltd.
                                                                                                                                                                                                                                                                                                             2006, 04:20:54 ;
                                                                                                                                                                                                                                                                                        Search time 26.4706 Seconds (without alignments)
49.601 Million cell updates/sec
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                                                                                                                        RESULT 1
US-08-175-158A-2
Sequence 2, Application US/08175158A
Patent No. 5986067
Patent No. 5986067
GENERAL INFORMATION:
APPLICANT: FUNK, Walter D.
APPLICANT: MACGILLIVRAY, ROSS T.A.
APPLICANT: MACGILLIVRAY, ROSS T.A.
APPLICANT: MOCOMORTH, ROBERT C.
TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSE: LAHIVE & COCKFIELD
                                                                                                                                                                                                US-08-683-262B-43
US-08-683-262B-45
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US-08-683-262B-46
US-08-683-262B-48
US-08-480-173A-48
US-08-480-1707-39
US-09-361-707-42
US-09-361-707-43
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Total number of Searched: Scoring table: Sequence: Title: Perfect score:

US-10-612-162A-3 74

SKLSMGSGLNLSEPN

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110 110 110 110 110 111 110 110 110 110	Result	Database
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CITY: Boston STATE: Massachusetts COUNTRY: USA ZIP: 02109 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII text CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/175,158A FILING DATE: 28-DEC-993 CLASSIFICATION DATA: APPLICATION NUMBER: US 07/832,029 FILING DATE: 06-FEB-1992 ATTORNEY/AGENT INFORMATION: NAME: DeConti, Giulio A. REGISTRATION NUMBER: UVI-005CP2 FILEDROMEY/AGENT INFORMATION: REFERENCE/DOCKET NUMBER: UVI-005CP2 TELEPAX: (617) 227-5941 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 698 amino acids TYPE: amino acid TYPE: amino acid TYPE: amino acid TYPE: amino acid TYPE: molecular TYPE: protein US-08-175-158A-2 Query Match Best Local Similarity 85.7%; pred. No. 0.022; Matches 12; Conservative 0; Mismatches 0;	; ADDRESSEE: LAHIVE & COCKFIELD ; STREET: 60 State Street, suite 510

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515 KLCMGSGLNLCEPN 528

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KLSMGSGLNLSEPN

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                                                                                                                                        Sequence 2, Application US/08470058 Patent No. 5817789
                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
TITLE OF INVENTION: Chimeric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 227-5941
[NFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 698 amino acids
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: DeConti, Giulio A.
REGISTRATION NUMBER: 31,503
REFERENCE/DOCKET NUMBER: UVI-005CP2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: WOODWORTH, ROBERT C.
TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/439,740 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                     12;
                                                                                                                                                                                                                                                                                         2 KLSMGSGLNLSEPN 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
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                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Floppy disk
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85.7%;
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                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                    Score 60; DB 2;
Pred. No. 0.022;
                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                     2; Indels
                                                                                                                                                                                                                                                                                                                                                                      Length 698;
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Best Local Similarity
Matches 12; Conserve
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SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amin-
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Patent No.
                                                                                                                                        COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FRASESEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
PILING DATE: 02-MAR-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
PILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Ph.D., J.D., Ja
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 0723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Heartlein, Michael W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: TKT9
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: CH
TITLE OF INVENTION: IN
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,058
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                          CITY: Boston
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STRANDEDNESS: unl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
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Concino, Michael F.
VENTION: CHIMERIC PROTEINS FOR USE
VENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
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85.7%;
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   07236/009002
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Pred. No.
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                                    Janis K.
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                                                                             Query Match
Best Local Similarity
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Best Local Similarity
Matches 12; Conserv
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino aci
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APPLICATION NUMBER: US/09/285,310
FILING DATE:
PRIOR APPLICATION DATA:
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TOPOLOGY: unknown
MOLECULE TYPE: protein
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LENGTH: 1074 amino acids
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                    NAME: Fraser, Ph.D., J.D., REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                       TELEPHONE:
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 891 KLCMGSGLNLCEPN 904
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                              KLSMGSGLNLSEPN 15
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                                                               Conservative
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85.7%;
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                                                                             Score 60; D
Pred. No. 0.
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Pred. No. 0.036;
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                                                                             DB 2; Length 1074; 0.036;
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APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE IN TRANSPORT
TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS
FILE REFERENCE: 10278-025004
CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1995-06-06
PRIOR FILING DATE: 1998-03-09
PRIOR FILING DATE: 1998-03-09
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US-09-753-385-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 5817789
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                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,058
FILLING DATE: 06-JUN-1995
CLASSIFICATION: 530
CLASSIFICATION: 530
                     TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/285,310 PRIOR FILING DATE: 1999-04-02
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TYPE: PRT
                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: TK:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey Fr
TITLE OF INVENTION: Chineric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS: ADDRESSEE: Hamilton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
SEQUENCE CHARACTERISTICS
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Local Similarity 85.7%;
les 12; Conservativo
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ZIP: 02173
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                                                                                                            TKT93-01
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Pred. No. 0.036;
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RESULT 9
US-09-285-310-4
; Sequence 4, Application US/09285310
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Best Local Similarity
""" ches 12; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION OF ANALYSIS
FILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRABER Ph.D., J.D., Janis K.
REGISTRATION NUMBER: 07236/009002
REFERENCE/DOCKET NUMBER: 07236/009002
                                                                                                                                                    Matches
                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                      TELEX: 200154
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 1410 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lemontt, Jeffrey F.
APPLICANT: CONCINO, MICHAEL F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617/542-8906
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STRANDEDNESS: un
TOPOLOGY: unknow
                                                                                                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                    Local Similarity es 12; Conserv
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STREET: 225 Franklin Street
                                                                                     1227
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                                                                                     KLCMGSGLNLCEPN 1240
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                                                                                                                                                                      Score 60;
Pred. No.
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Pred. No. 0.05;
                                                                                                                                                      Mismatches
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                                                                                                                                                                                      DB 2;
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APPLICANT: Heartlein, Michael W.
APPLICANT: Hemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
APPLICANT: Concino, Michael F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS
FILE REFERENCE: 10278-025004
CURRENT APPLICATION NUMBER: US/09/753,385
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 08/470,058
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR FILING DATE: 1998-03-09
PRIOR FILING DATE: 1998-03-09
PRIOR APPLICATION NUMBER: US 09/037,188
PRIOR APPLICATION NUMBER: US 09/285,310
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NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09753385
Patent No. 6858578
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SEQUENCE CHARACTERISTICS:
LENGTH: 1410 amin
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APPLICANT: Heartle
APPLICANT: Lemontt
APPLICANT: Concino
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                             -09-753-385-4
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/037,18:
APPLICATION NUMBER: 09/037,18:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: FYRBER; Ph.D., J.D., Ja:
NAME: FYRBER; Ph.D., J.D., Ja:
REGISTRATION NUMBER: 34,819
REGISTRENCE/DOCKET NUMBER: 0723
TRLECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                        PRIOR FILING DATE: 1999-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: unknown MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOCTWARE: PastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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ZIP: 02110-2804
COMPUTER READABLE FORM:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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CONCINO, Michael F.
NVENTION: CHIMERIC PROTEINS FOR USE
NVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       225 Franklin Street
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Pred. No. 0.05;
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FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 15968
LENGTH: 439
TYPE: PRT
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                                 Query Match
Best Local Similarity
Watches 8; Conserve
                                                                                                                                                                                         ; NAME/KEY: UNSURE
; LOCATION: (12)
; OTHER INFORMATION:
US-09-248-796A-15968
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US-09-248-796A-15968
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Candida albicans
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APPLICATION NUMBER: 827,313
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/7230
FILING DATE: 27-JAN-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 696
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SKLSMGSGLNLSEP 14
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                                                                Conservative
                                                                                                                                                                                                                           Identity of amino acid sequences at the above locations are unknown
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85.7%;
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57.1%;
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                                                                                           Score 46; DB 2;
Pred. No. 4.5;
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Pred. No. 0.62;
0; Mismatches
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                                                                   Mismatches
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                                                                                                                             Length 439;
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PRIOR APPLICATION NUMBER: PCT/JP98/01783;
PRIOR FILING DATE: 1998-04-17
PRIOR APPLICATION NUMBER: JP 9/310027
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: JP 9/116570
PRIOR FILING DATE: 1997-04-18
NUMBER OF SEQ ID NOS: 73
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 72
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; ORGANISM: Homo sapiens
US-09-839-479-71
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US-09-839-479-71
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                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 71
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 71, Application US/09839479
Patent No. 6727222
GENERAL INFORMATION:
APPLICANT: Jones, Michael H.
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 06501-042002
                                                                  Matches
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Patent No. 6596482
                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 06501-042002
CURRENT APPLICATION NUMBER: US/09/839,479
CURRENT FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: US 09/418,710
PRIOR FILING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: PCT/JP98/01783
PRIOR FILING DATE: 1998-04-17
PRIOR FILING DATE: 1998-04-17
                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: JP 9/310027
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: JP 9/116570
PRIOR FILING DATE: 1997-04-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/418,710 CURRENT FILING DATE: 1999-10-15
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TYPE: PRT
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SLGSGLGLSEGN 1465
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                                                                  Conservative
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75.0%;
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75.0%;
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Pred. No.
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Pred. No. 91;
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RESULT 15

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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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length: 0
length: 2000000000
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74
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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Listing first 45 summaries
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             US-10-612-162-3
US-10-378-094-15
US-10-378-094-15
US-10-38-1-266-745-5
US-10-266-745-5
US-10-266-745-5
US-10-384-060-3
US-10-314-494-3
US-10-429-497-4
US-10-429-535-4
US-10-429-653-4
US-10-429-653-4
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US-10-429-661-4
US-10-429-661-4
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US-10-429-653-4
US-10-429-653-4
US-10-429-654-4
US-10-378-0942-6
US-10-378-0942-6
US-10-378-0942-6
US-10-378-0942-6
US-10-378-0942-6
US-10-384-060-2
US-10-384-060-2
US-10-383-201-14
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RESULT 1 US-10-612-162- ; Sequence 3; ; Dublication ; GENERAL INFO ; APPLICANT: ; TITLE OF IN		45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	
SULT 1 :-10-612-162-3 :-10-612-162-3 :-uence 3, Application US/10612162 Publication No. US20040014145A1 GENERAL INFORMATION: APPLICANT: Dade Behring Marburg Gm TITLE OF INVENTION: Carbohydrate d			43 58	43 58		58	63	74	74	74	74	74	77	81	81	60 81		60 81.1		
cation US/: S200400141 ON: Behring Mai		.1 1972	Ļ	.1 1969	Ļ	Ļ	Մ	ω	W	'n	'n			۲.	.1 1410			.1 698		
ULT 1 10-612-162-3 lequence 3, Application US/10612162 ublication No. US20040014145A1 ENERAL INFORMATION: APPLICANT: Dade Behring Marburg GmbH APPLICANT: Dade Behring Marburg GmbH	ALIGNMENTS	3 US-09-839-479-21	Sn	4 US-10-376-537-72		3 US-09-864-761-36754		-	-	-	-	5 US-10-769-514-10	-	-		SU	ŭ	6 US-11-177-506-46	5 US-10-513-523-3	
n (CDT)-specific			Sequence 71, Appl	72,		Seguence 36754, A				180		10,		Sequence 804, App	Sequence 4, Appli			æ	Sequence 3, Appli	

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TITLE OF INVENTION: Carbohydrate delicient transferrint TITLE OF INVENTION: antibodies, their preparation and FILE REFERENCE: 2002/B001
CURRENT APPLICATION NUMBER: US/10/612,162
CURRENT FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 10230550.1
PRIOR FILING DATE: 2002-07-05
PRIOR FILING DATE: 2003-05-19
PRIOR FILING DATE: 2003-05-19
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3
APPLICANT: PRIOR, Christopher P.
APPLICANT: LAI, Char-Huei
APPLICANT: LAI, Char-Huei
APPLICANT: TURNER, Homeyoun
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN FUSION PROTEINS
FILE REFERENCE: 54710-5001-01-US
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT FILING DATE: 2003-03-04
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/334,059
                                                                                                                                                                                                                                                                                                                     US-10-378-094-15
; Sequence 15, Application US/10378094
; Publication No. US20030221201A1
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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RESULT 3
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; NAME/KEY: misc feature
; OTHER INFORMATION: C1 subdomain of transferrin
US-10-378-094-15
                                                                                                                                                                                                                                           US-09-891-126-5
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CURRENT APPLICATION NUMBER: US/10/384,060
CURRENT APPLICATION NUMBER: US/10/331,494
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 80
Sequence 5, Application US/09891126
Patent No. US2002072596A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Transferrin Polynucleotides, Polypeptides, and Antibodies
FILE REFERENCE: PT035P1
CURRENT APPLICATION NUMBER: US/09/891,126
CURRENT FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/171,595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 36, Application US/10384060 Publication No. US20030226155A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PA
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NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn version 3.2
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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APPLICANT: PRIOR, Christopher P.
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 42
TYPE: PRT
ORGANISM: Homo sapiens
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PRIOR APPLICATION NUMBER: US/09/891,126
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR APPLICATION NUMBER: FOT/US00/34769
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR FILING DATE: 1999-12-23
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 5
LENGTH: 328
TYPE: PRT
ORGANISM: Homo sapiens
US-10-266-745-5
                                              APPLICANT: PRIOR, Christopher P.
APPLICANT: LAI, Char-Huei
APPLICANT: SADEGHI, Homayoun
APPLICANT: SADEGHI, Homayoun
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN FUSION PROTEINS
FILE REFERENCE: 54710-5001-01-US
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT FILING DATE: 2003-03-04
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR FILING DATE: 2002-08-30
PRIOR FILING DATE: 2001-11-30
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2001-08-30
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-891-126-5
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NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/10378094
Publication No. US20030221201A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/10266745 Publication No. US20030149256A1 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 5
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Best Local
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TITLE OF INVENTION: Transferrin Polynucleotides, Polypeptides, and Antibodies
FILE REFERENCE: PT035P1
CURRENT APPLICATION NUMBER: US/10/266,745
CURRENT FILING DATE: 2002-10-09
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Pred. No.
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Sequence 3, Application US/10231494
Publication No. US20040023334A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
CURRENT APPLICATION UNMBER: US/10/231,494
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2001-11-30
NUMBER OF SCR OT NOCE 30
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; FEATURE:
; NAME/KEY: MISC_FEATURE
; OTHER INFORMATION: Mature Transferrin Protein
US-10-378-094-3
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CURRENT APPLICATION NUMBER: US/10/384,060
CURRENT FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: 2002-08-30
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Best Local Similarity 85.7%;
Matches 12; Conservative
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APPLICANT: PRIOR, Christopher P.
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 80
SOFTWARE: PatentIn version 3.2
NUMBER OF SEQ ID NOS: 30
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NAME/KBY: MISC_FEATURE
OTHER INFORMATION: Mature Transferrin Protein
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TYPE: PRT
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Local Similarity 85.7%;
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Pred. No. 0.079;
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Pred. No. 0.079;
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Sequence 4, Application US/10429497

Publication No. US20040219098A1

GENERAL INFORMATION:

APPLICANT: Sandhu, Jasbir

TITLE OF INVENTION: Methods for The Treatment of Tumors

FILE REFERENCE: 2537.000006

CURRENT APPLICATION NUMBER: US/10/429,497

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 6

SOFTWARE: Patentin version 3.1

SEQ ID NO 4
                                                                                                                     ; LENGTH: 679
; TYPE: PRT
; ORGANISM: HOMO &
US-10-429-497-4
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-429-482-4
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US-10-429-497-4
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; LENGTH: 679
; TYPE: PRT
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TITLE OF INVENTION: Composition Useful For The Diagnosis, Imaging and Treatment Of
FILE REFERENCE: 2537.000001
CURRENT APPLICATION NUMBER: US/10/429,482
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
                                                   Query Match
Best Local Similarity
Matches 12; Conserv
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Local Similarity 85.7%;
nes 12; Conservative
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KLCMGSGLNLCEPN 509
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85.7%;
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Pred. No. 0.079;
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Pred. No. 0.079;
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                                                             Mismatches
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RESULT 11 US-10-429-515-4

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Sandhu, Jasbir

TITLE OF INVENTION: Composition Useful For Treatment Of Tumors

FILE REFERENCE: 2537.000007

CURRENT APPLICATION NUMBER: US/10/429,635

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 8

SOFTWARE: Patentin version 3.1

SEQ ID NO 4

LENGTH: 679

TYPE: PRT

ORGANISM: Homo sapiens

US-10-429-635-4
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ITITLE OF INVENTION: Methods For The Treatment Of Tumors

FILE REFERENCE: 2537.000005

CURRENT APPLICATION NUMBER: US/10/429,515

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 6

SOFTWARE: Patentin version 3.1

SEQ ID NO 4

LENGTH: 679

TYPE: PAT

ORGANISM: Homo sapiens

US-10-429-515-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The Treatment of Tumors
FULE REFERENCE: 2537.000003
CURRENT APPLICATION UMMER: US/10/429,598
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
US-10-429-598-4
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US-10-429-635-4
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Query Match 81.1%;
Best Local Similarity 85.7%;
Matches 12; Conservative
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Publication No. US20040219100A1
GENERAL INFORMATION:
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Best Local Similarity 85.7%;
Matches 12; Conservative
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Best Local Similarity 85.7%;
Matches 12; Conservative
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 Score 60; DB 5;
Pred. No. 0.079;
0; Mismatches
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Pred. No. 0.079;
0; Mismatches
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Search completed: June 13, 2006, 04:56:23
Job time : 86.25 secs
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TITLE OF INVENTION: Methods Useful For The Diagnosis, Imaging and Treatment Of
FILE REFERENCE: 2537.00004
CURRENT APPLICATION NUMBER: US/10/429,659
CURRENT FILING DATE: 2003-04-30
NUMBER OF SEQ ID DOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/10429659
Publication No. US20040219103A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Compositions For Drug Delivery
FILE REFERENCE: 2537,00009
CURRENT APPLICATION NUMBER: US/10/429,653
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
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TYPE: PRT
ORGANISM: Homo sapiens
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Title:

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length: 2000000000
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Match
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1: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US09_NEW_PUB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US11_NEW_PUB.pep:*

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8: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
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Listing first 45 summaries
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IS-10-953-349-3255

IS-11-293-697-2635

IS-10-953-349-29403

IS-10-953-349-29403

IS-10-953-349-15250

IS-10-953-349-15249

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IS-10-953-349-13180

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IS-10-953-49-16374

IS-10-953-349-2063

IS-10-953-349-13573

IS-10-953-349-13572

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Sequence 3, Appli
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Sequence 18574, A
Sequence 20639, A
Sequence 13572, A
Sequence 13572, A
Sequence 20637, A
Sequence 2637, A
Sequence 3265, Ap
Sequence 3265, Ap
Sequence 29404, A
Sequence 29404, A
Sequence 29404, A
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Sequence 15250, A
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Sequence 15249, A
Sequence 13119, A
Sequence 34845, A
Sequence 34845, A
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Publication No. US20060105387A1

GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
APPLICANT: Prior, Christopher P.
APPLICANT: Sadeghi, Homosyoun

ITITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
CURRENT APPLICATION NUMBER: US/10/515,429

CURRENT FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 00/485,404
PRIOR APPLICATION NUMBER: US 00/485,404
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
RESULT 2
US-10-515-429-2
; Sequence 2, Application US/10515429
; Publication No. US20060105387A1
; GENERAL INFORMATION:
; APPLICANT: Prior, Christopher P.
; APPLICANT: Turner, Andrew J.
; APPLICANT: Sadeghi, Homayoun
; TITLE OF INVENTION: Transferrin Fusion Protein Libraries
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; OTHER INFORMATION: Mature transferrin protein
US-10-515-429-3
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US-10-515-429-3
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Matches 12; Conserv
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US-10-953-349-18594
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Minimum DB Maximum DB

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FILE REFERENCE: 054710-5007-WO

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; TYPE: PRT; ORGANISM: Glycine max US-10-953-349-18406
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; Sequence 18406, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
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US-10-953-349-18405
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CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin Ver. 2.1
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 18406
LENGTH: 128
                                     NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 18405
LENGTH: 167
                                                                                                                                                                                                                                        Sequence 18405, Application US/10953349 Publication No. US20060107345A1 GENERAL INFORMATION:
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                                                                                                               APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE DETERMINED DI
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
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TITLE OF INVENTION: SEQUENCE-DETERMINED
TITLE OF INVENTION: ENCONDED THERBY
TYPE: PRT ORGANISM: Glycine max
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SKLAVGDGVPVSQP 37
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85.7%;
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APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 18404
LENGTH: 188
TYPE: PRT
ORGANISM: Glycine max
US-10-953-349-18404
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                                PRIOR APPLICATION NUMBER: US 60/404,249
PRIOR FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 60/396,594
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: US 60/387,292
PRIOR FILING DATE: 2002-06-07
PRIOR FILING DATE: 2002-06-07
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Publication No. US20060107345A1
GENERAL INFORMATION:
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Best Local
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APPLICANT: Hakes, David
APPLICANT: Chen, Xi
APPLICANT: Chen, Xi
APPLICANT: Bowe, Caryne
TITLE OF INVENTION: ERYTHROPOIETIN: REMODELING AND
TITLE OF INVENTION: GLYCOCONJUGATION OF ERYTHROPOIETIN
FILE REFERENCE: 040853-01-5083-US02
CURRENT APPLICATION NUMBER: US/11/183,218
CURRENT FILING DATE: 2005-07-15
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PRIOR FILING DATE: 2003-04-09
PRIOR APPLICATION NUMBER: PCT/US02/32263
PRIOR FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US 60/407,527
PRIOR FILING DATE: 2002-08-28
PRIOR FILING DATE: 2002-08-28
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APPLICATION NUMBER: US 60/344,692 FILING DATE: 2001-11-19
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US-10-953-349-20639
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US-10-953-349-13574
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US-11-183-218-46
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Publication No. US20060107345A1
GENERAL INFORMATION:
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Publication No. US20066107345A1

GENERAL INFORMATION:

APPLICANT: ALEXANDROV, Nickolai et al.

TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES

TITLE OF INVENTION: ENCONDED THERBY

FILE REFERENCE: 2750-1579PUS2

CURRENT APPLICATION NUMBER: US/10/953,349

CURRENT FILING DATE: 2004-09-30
                                                                                                                                                      NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 20639
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 46
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LENGTH: 409
                               Matches
                                            Query Match
Best Local Similarity
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
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SOFTWARE: PatentIn version 3.3
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TYPE: PRT
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Pred. No. 9
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US-10-953-349-13572
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                                                                                                                                                                       Sequence 13572, Application US/10953349 Publication No. US20060107345A1 GENERAL INFORMATION:
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APPLICANT: ALEXANDROV, NICKOLAI et al.
APPLICANT: ALEXANDROV, NICKOLAI et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 2638
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT FILLING DATE: 2004-09-30
CURRENT FILLING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 11573
LENGTH: 424
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Publication No. US20060107345A1
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                APPLICANT: ALEXANDROY, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DI
TITLE OF INVENTION: SECONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILLING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
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ORGANISM: Glycine max
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; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1). (290)
; OTHER INFORMATION: hypothetical protein
US-10-471-571A-180
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US-10-471-571A-180
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US-10-953-349-20637
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Best Local Similarity
7; Conservation
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                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: GB-0107661.1
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 5642
SOFTWARE: SegWin99, version 1.03
SEQ ID NO 180
LENGTH: 290
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 20637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20637, Application US/10953349
publication No. US20066107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: NICKOLAI et al.
APPLICANT: SEQUENCE TERRMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
FILE REFERENCE: 2750-1579PUS2
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LENGTH: 472
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Publication No. US20060115490A1
GENERAL INFORMATION:
APPLICANT: CHIRON SpA
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FILE REFERENCE: P026927WO
CURRENT APPLICATION NUMBER: US/10/471,571A
CURRENT FILING DATE: 2003-09-12
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TYPE: PRT
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TYPE: PRT
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Local Similarity 46.2%;
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424 IQLGRGLDIGEPN 436
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1 SKLSMGSGLNLSEPN 15
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Pred. No. 11;
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9.2;
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CURRENT APPLICATION NUMBER: US/11/293,697
CURRENT FILING DATE: 2005-12-05
PRIOR APPLICATION NUMBER: US/10/108,260
PRIOR FILING DATE: 2002-03-28
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4653
LENGTH: 208
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Search completed: June 13, 2006, 04:56:53 Job time: 6.07353 secs
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PATENTIN Version 3.3
SEQ ID NO 3265
LENGTH: 275
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                 Sequence 3265, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
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APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: Novel full length cDNA
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This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSWGKEDLIWELL, TYEDSIAKIMNGRADAMSLDGGF, SKLSMGSGLMISSEN and YEKYLGEEYWAN (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for
                                                                                                                                                                                                                                                                                                                                      New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating
                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 3; 21pp; German.
                                                                                                                                                                                                                                                                                                                   need for immobilization.
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Aay54711 Human sub	Aay54710 Human sub	Adv70479 Moraxella	Abr31410 Human can	Abr30754 Human can	Abr38454 Human can	Abr38289 Human can	Abr38381 Human can	Abr31578 Human can	Abr30689 Human can	Adw93469 SARS coro	Adl21451 125P5C8 p	Abr30870 Human can	Abr30379 Human can	Abr31569 Human can	Abr30664 Human can	Abr30735 Human can	Abr31106 Human can	Abr31310 Human can	8	Adz07250 Hepatitis	Adz06623 Hepatitis	

ALIGNMENTS

antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. Human CDT peptide fragment EP1378521-A1. ADG46077; ADG46077 standard; peptide; 05-JUL-2002; 2002DE-01030550 19-MAY-2003; 2003EP-00011334. 07-JAN-2004. Homo sapiens. 25-MAR-2004 (DADE-) DADE BEHRING MARBURG GMBH (first entry) #3 15 B

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                                                                                           The present invention describes an assay for a protein having at least two isoforms with different glycosylation patterns. The assay involves contacting a sample containing the protein with a proteolytic enzyme, followed by detecting the content or relative content of at least one peptide fragment produced by proteolysis. Also described is a kit for the assay method comprising the proteolytic enzyme and a substrate bound specific binding partner (S1) for at least 2 of the isoforms of the proteins. The method can be used for assaying isoforms of proteins according to their glycosylation pattern to determine their concentration or relative concentration in the sample or material (e.g. blood). The method avoids use of antibodies for distinguishing between glycosylated isoforms of the proteins. The present sequence represents a transferrin peptide fragment which is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 15; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Assay for differentiating protein isoforms to determine their concentrations in sample e.g. blood, involves contacting the sample with proteolytic enzyme to produce peptide fragment by proteolysis followed by
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ADS13393 standard; peptide; 15

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                                                                                                                                                                                                                                                       Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated marker for rheumatoid arthritis selected from one of many (around 400) markers defined in the specification. Rheumatoid arthritis is a chronic inflammatory disorder of the small joints which is estimated to affect 2.1 million people in the
                                                                                                                                                                                                                                                                                          Human rheumatoid arthritis marker peptide - SEQ ID 180.
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               uptake.
                             Transferrin-binding molecules useful for eliciting antibodies to bacterial transferrin
                                                                           WPI; 2005-038740/04
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                                                                                                                                                                                                                                                           US2004258695-A1
                                                                                                                                                                                                                                                                                                                     neuroprotective;
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                                                                                                                                                                                                                                                                                          The invention comprises a molecule (e.g. peptide) which is capable of: binding to a region of a transferrin protein that is recognized by a bacterial transferrin binding protein; and eliciting an antibody to the bacterial transferrin binding protein. The transferrin binding molecule of the invention is useful for preventing and treating bacterial infections (e.g. bacterial meningitis and otitis media). The present amino acid sequence represents a peptide that is bound by transferrin
                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transferrin-binding molecules useful for eliciting antibodies to bacterial transferrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-038740/04.
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                                                                                                                                                                                                                                                                binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-MAR-2005
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                                                                                                                Similarity
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                      LSMGSGLNLSEPN 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 42; 27pp; English.
                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                             74.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibacterial; bacterial meningitis; antibacterial; tis media; auditory; transferrin binding protein B.
                                                                                Score 55; DB
Pred. No. 0.01
0; Mismatches
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Pred. No. 0.01;
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                                                                                                                                             Length 15;
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                                                                                                                                                                                                                                                                                                                The invention relates to a novel method for analyzing a mixture of proteins. The method involves selecting sets of peptides from several peptides, sequencing one or more portions of the peptides in two or more sets of peptides, to produce a data set, and determining the likelihood of the presence of two or more proteins in the mixture of proteins based on two or more data sets. The invention further comprises an article of manufacture comprising a computer-readable media with computer-readable instructions for carrying out the method. The method is useful for analyzing a mixture of proteins. The method is useful for seriening and characterization of protein mixtures. This sequence represents a human serum albumin (HSA) peptide fragment used in the novel protein analysis
                                           10-MAR-2005
                                                                     ADV23202;
                                                                                                                                                                                                                                                                               Sequence 15 AA;
                                                                                                                                                                                                                                                                                                         method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; SEQ ID NO 82; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analyzing mixture of proteins by sequencing portions of peptides in two or more sets of peptides from several peptides, to produce data set, determining likelihood of presence of proteins in mixture of proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-162781/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Regnier FE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-2003; 2003US-0490149P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein purification; chromatography; screening
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                                                                                               ADV23202 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    based on data sets.
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                                           (first entry)
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                                                                                                                                                                                                                                      74.3%;
84.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhang X;
                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                      Score 55;
Pred. No.
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                                                                                                                                                                                                                                                   9;
                                                                                                                                                                                                                                                     Length 15;
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HBV immunogenic peptide #3

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cc peptides above and thus presents the peptides, a population of such cells for modulating an immune response to a polypeptide of interest, a composition at least one set of the peptides (and a carrier and/or comprising at least one set of the peptides (and a carrier and/or comprising at least one set of the peptides (and a carrier and/or comprising at least one set of the peptides (and a carrier and/or comprising administering to a patient in need at least one set of the peptides (and a carrier and/or condition associated with the presence of a polypeptide of interest comprising administering to a patient in need at least one set of the peptides, a method for treatment and/or prophylaxis of a disease or condition associated with the presence of a polypeptide of interest is also a composition of matter for modulating an immune response in a composition of matter for modulating an immune response in a composition of matter for modulating an immune response in a composition of matter for modulating an immune response in a composition organism or a cancer, and produced by a pathogenic organism or a cancer, and produced by a pathogenic organism or a cancer, and produced by a pathogenic organism or a cancer, and produced by a produced by a cancer selected from melanoma, lung cancer, croacer, stomach cancer, prostate cancer, colon cancer, pancreatic cancer, stomach cancer, bladder cancer, kidney cancer, post transplant composition of a medicament for the treatment of a disease or condition is associated with the presence or aberrant expression of a target antigen, where the antigen-presenting colls or their precursors are useful in the antigen-presenting colls or their precursors are useful in the antigen to express a processed or modified form of the antigen to express a processed or modified form of the antigen to express a processed or modified form of the antigen to overlapping immunogenic peptides derived from a Hepatitiis of the present sequence is one of a set of overlapping immunogenic peptides derived from 
Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        individual peptides of a respective set comprise different portions of an amino acid sequence corresponding to a single polypeptide of interest and display partial sequence identity or similarity to at least one other peptide of the same set of peptides (i.e. they are overlapping). Also included are an antigen-presenting cell which has been contacted with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vaccine; virucide; antigen; autoimmune disease; infection; immune modulation; cancer; neoplane; prostatic; melanoma; lung tumor; breast tumor; uterine cervix tumor; prostatic cancer; colon tumor; pancreas tumor; stomach tumor; bladder tumor; kidney tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of at least one set of peptides in the preparation of a medicament for modulating an immune response, and for treating cancer or yeast, viral, bacterial, protozoal and mycoplasma infections.
                                                                                                             Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preparation of a medicament for modulating an immune response,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 1622; 645pp; English.
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25-MAR-2004; 2004AU-00901589.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to the use of at least one set of peptides in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kent
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                 Conservative
                                       47.3%;
                 Score 35; DB Pred. No. 41; 0; Mismatches
                                  DB 41;
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6 GSGLNLSEPN 15

HBV immunogenic peptide #2

10-MAR-2005

(first entry)

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RESULT 11
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human prothrombin kringle-1 and kringle-2 polypeptides that inhibit angiogenesis, useful for treating cancer.
                                                     ADV23201
                                                                                                                                                                                                                                                                     Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        skin; laryngeal; uterine; colon; lung; bone marrow; cytostatic;
anti-angiogenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human prothrombin kringle-2; prothrombin; kringle domain; cancer therapy; anti-endothelial cell proliferation; anti-tumour; angiogenesis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY70486;
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                                                                              ADV23201 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 7; Page 19; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rhim TY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthesised peptide-1 from human prothrombin kringle-2 protein.
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                                                                                                                                                                                SMGSGLNLSEP 14
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                                                                                                                                                                                                              Conservative
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Competition relates to the use of at least one set of peptides in the individual peptides of a medicament for modulating an immune response, where a minimal acid sequence corresponding to a single polypeptide of interest and competite of the same set of peptides (i.e. they are overlapping). Also included are an antigen-presenting cell which has been contacted with the competites for modulating an immune response to a polypeptide of interest, a competite for modulating an immune response to a polypeptide of interest, a competition for modulating an immune response to a polypeptide of interest, a competition for modulating an immune response to a polypeptide of interest, a composition comprising at least one set of the peptides (and a carrier and/or compising at least one set of the peptides (and a carrier and/or composition of matter for modulating an immune response to a polypeptide of interest cof the peptides, a method for treatment and/or prophylaxis of a disease condition associated with the presence of a polypeptide of interest cof and a composition of matter for modulating an immune response to a polypeptide of interest can accordance or condition associated with the presence of a polypeptide of interest can accordance or condition associated polypeptide that is a polypeptide cof protozoans and mycoplasmas. The polypeptide that is a polypeptide composition or a pathogenic organism or a cancer, and produced by a pathogenic organism or a cancer and produced by a pathogenic organism or a cancer, occur, post transplant cof protozoans and mycoplasmas. The disease ro condition associated cancer, bladder cancer, bladder cancer, bladder cancer, because, post transplant cof a medicament for the treatment of a disease or condition is associated with the presencing cells or their precursors are useful in the presencing or a medicament for the treatment of a disease or condition is associated with the presence of the target antigen to be protozoans and mycoplasmas protosessed or modified form of the antigen to organi
Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                              Sequence
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25-MAR-2004;
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Score 33; DB
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibody production against allergenic antigen (Ag) is specifically suppressed by treatment with the junctional segment of the alpha and/or beta chain of the Ag receptor of suppressor T (Ts) cells (TCR) induced by tolerogenic Ag-PBG conjugates. Induction of antigen-specific suppression by treatment of mice with TCR alpha-chain peptides was demonstrated using the peptides given in AAR62912-17. The peptides of AAR62914 corresponds to the J-alpha region of the CDR3 segment of the alpha chain of ovalbumin-specific Ts cells. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complementarity determining region; immunosupprantibody engineering; suppressor T-lymphocyte; monoclonal antibody; myeloma; IgG.
                                                            Peptide from library spanning whole of hsp65.
                                                                                                                           11-NOV-1996
                                                                                                                                                                                                                                             AAR94866 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic peptide(s) for effecting immuno:therapy - have an amino acid sequence comprising a portion of the CDR3 region of a I-cell receptor.
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Hsp; heat shock protein; Mycobacterium bovis; microbial; diagnosis;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR94773-R94878 are overlapping peptides of a library spanning the whole of microbial hsp65 (heat shock protein 65). Eleven antigens were prepared from the library and used to immunise mice, after 10 days the mice received injections of pristane to induce arthritis and the animals examined for incidence of arthritis. A preferred anti-arthritic peptide was found to correspond to residues 21-31 of full length hsp65 (see AAR94772). Peptides conty. this sequence are used in the diagnosis, treatment and prophylaxis (may be used in a vaccine) of autoimmune diseases such as rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rulypeptide derived from bacterial heat shock protein 65 - f diagnosis, prophylaxis and treatment of auto-immune disease rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                   Heat shock protein; hsp58; autoimmune disease; hsp65; vaccine; rheumatoid arthritis; antigen; infectious disease; prophylacti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-2003
08-APR-1998
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                                                                                                                                                          03-APR-1997.
                                                                                                                                                                                                                                                                                        Synthetic.
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                                                    27-SEP-1995;
                                                                                                     26-SEP-1996;
                                                                                                                                                                                                              WO9711966-A1
                                                                                                                                                                                                                                                                Mycobacteriaceae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mycobacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW43544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW43544 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Fig 1; 88pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thompson SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PEPT-)
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(PEPT-) PEPTIDE THERAPEUTIC LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6
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                                                                                                                                                                                                                                                                                                                                             arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                hsp68 derived peptide (group 10 #4).
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                                                                                                                                                                                                                                                                                                                                                                   antigen; infectious disease; prophylactic;
                                                                                                                                                                                                                                                                                                                                               PIA.
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Pred. No. 3.3e+02;
1; Mismatches 1
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RESULT 15
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This peptide is one of a library (see AAW43451-W43556) which represents the mycobacterial heat shock protein, hsp65 and is designed to study a novel method for the treatment of autoimmune disease e.g. rheumatoid arthritis. This fragment has been allocated as a group 10 peptide which is composed of the fragments found in AAW43541 to AAW43550. Mycobacterial hsp65 is known to be an immunodominant antigen in a number of infectious diseases and is linked to pristane induced arthritis (PIA) in vitro, however heat shock proteins and peptides derived from microbial sources may act as self antigens and thus have limited clinical use. The human hsp65 homologue, hsp55, or fragments of the hsp58 protein (see AAW14948-W14950) may be useful in the development of vaccines for prophylaxis or treatment of an autoimmune disease such as rheumatoid arthritis. (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                Sette A,
Celis E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                 Vaccine composition for treating or preventing hepatitis B vinfection, and/or for stimulating an immune response to HBV, HBV peptide epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                       14-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-212851/19.
                                                                                                        WPI; 2002-643192/69.
                                                                                                                                                                         (SETT/)
                                                                                                                                                                                                               08-SEP-2000; 2000WO-US024802
                                                                                                                                                                                                                                          08-SEP-2000; 2000WO-US024802
                                                                                                                                                                                                                                                                    14-MAR-2002.
                                                                                                                                                                                                                                                                                               WO200219986-A1.
                                                                                                                                                                                                                                                                                                                         Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                   virucide; hepatotropic;
                                                                                                                                                                                                                                                                                                                                                                 Hepatitis B virus; HBV;
                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis B virus epitope #3179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABJ08961 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 1c; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polypeptide(s) derived from microbial heat shock protein - useful for treatment of autoimmune disease esp. arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson JS,
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                                                                                                                                                                       SETTE A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LSVGHGLN 11
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                                                                                                                              Sidney J,
Kubo RT,
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                                                                                                                                Southwood S, Vit
Grey HM, Chesnut
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                                                                                                                                                                                                                                                                                                                                                     epitope; vaccine; HBV infection; hepatitis; antiinflammatory.
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Pred. No. 3.3e+02;
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Disclosure; Page 179;

228pp; English

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The present invention relates to a composition comprising at least

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Post-processing: Minimum Match
Maximum Match
Listing first
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length:
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Match
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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S21238
                                                   A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-15 <TJO>
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C36198
                                                                                  A; Reference number: A; Accession: C36198
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                                                                                                                                                                                                                                                                                                                                                                         C; Genetics:
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                 Matches
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Best Local :
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Searched:

Scoring table: Sequence:

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ty meany charm no	Ic heavy chain D.I	milk band B protei	20K protein 5403 -	protein QA100039 -	prostaglandin D-sy	major fat-globule	T-cell receptor al	protein kinase (EC	glutathione-disulf	hexokinase (EC 2.7	malate dehydrogena	fructose-bisphosph	T-cell receptor al	T-cell receptor al	Ig heavy chain V r	collagen alpha 2(X

ALIGNMENTS

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T-cell receptor beta chain J region - chicken (fragment)
;Species: Gallus gallus (chicken)
C;Date: 28-Mar-1991 #sequence_revision 28-Mar-1991 #text_change 21-Jul-2000
C;Accession: C36198
R;Tjoelker, L.W.; Carlson, L.W.; Lee, K.; Lahti, J.; McCormack, W.T.; Leider
R;Tjoelker, L.W.; Carlson, L.W.; Tee, K.; Lahti, J.; McCormack, W.T.; Leider
Proc. Natl. Acad. Sci. U.S.A. 87, 7856-7860, 1990
A;Title: Evolutionary conservation of antigen recognition: the chicken T-cel
A;Reference number: A36198; MUID:91045896; PMID:2236002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Description: catalyzes the six-electron reduction of sulfite to sulfide A;Pathway: the terminal oxidase in the sulfate-reduction pathway C;Keywords: heterohexamer; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Pierik, A.J.; Duyvis, M.G.; van Helvoort, J.M.L.M.; Wolbert, R.B.G.; Hagen, W.R. Eur. J. Biochem. 205, 111-115, 1992
A;Title: The third subunit of desulfoviridin-type dissimilatory sulfite reductases A;Reference number: S21197; MUID:92209491; PMID:1555572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hydrogensulfite reductase (EC 1.8.99.3) beta chain - Desulfovibrio vulgaris (fragment) N;Alternate names: bisulfite reductase; desulfofuscidin; desulforubidin; desulfoviridin; C;Species: Desulfovibrio vulgaris
                                                                                                   A;Cross-references: UNIPARC:UPI000011511E; EMBL:M37799; NID:g212737; PIDN:AAA62754.1; PIII
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: dsvB
C;Complex: heterohexamer; two alpha, two beta and two gamma chains
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A;Experimental source: strain Hildenborough
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Score
Pred.
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Pred. No. 1.8e+03;
2; Mismatches 4
No.
DB 2;
1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.
                             Length 15;
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                                                                                                                                                                                                                                                                                                                                              receptor be
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C;Species: Streptococcus oralis
C;Date: 29-Nov-1995 #sequence_revision 01-Aug-1997 #text_change 31-Dec-2004
C;Accession: S49409; S38211
C;Accession: S49409; S38211, Garcia, E.; de la Campa, A.G.
R;Fenoll, A.; Munoz, R.; Garcia, E.; de la Campa, A.G.
Mol. Microbiol. 12, 587-598, 1994
A;Title: Molecular basis of the optochin-sensitive phenotype of pneumococcus:
A; Molecule type: DNA
A; Residues: 1-15 < FEN:
                                                          A;Reference number: S49398; MUID:95020593; A;Accession: S49409
                                                                                                                                                                                                                                              H+-transporting two-sector ATPase (EC 3.6.3.14) chain a -
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A; Residues: 1-15 < CAS>
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J. Exp. Med. 177, 811-820, 1993
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A;Title: p180, a novel recycling transmembrane glycoprotein with A;Reference number: A36315; MUID:90258846; PMID:2188094
A;Accession: A36315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Homo sapiens (man)
C;Date: 25-Jan-1991 #sequence_revision 25-Jan-1991 #text_change 30-Sep-1993
C;Accession: A36315
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                                A;Status: preliminary
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C;Species: Mus musculus (house mouse)
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Matches 5
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Pred. No. 1.8e+03;
0; Mismatches 2;
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Pred. No. 1.8e+03;
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A;Title: T cell receptor selection by and recognition of A;Reference number: PH1430; MUID:93171821; PMID:8436911 A;Accession: PH1442 A;Molecule type: mRNA A;Residues: 1-15 cCA2> A;Cross-references: UNIPARC:UPI00001769BB A;Cross-references: UNIPARC:UPI00001769BB A;Experimental source: Cytolytic T-lymphocyte C;Superfamily: immunoglobulin V region; immunoglobulin ho
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$26527

T-cell receptor alpha chain V region (clone Cw3/A8, A3/74.1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
A;Variety: clone Cw3/A8, A3/74.1
C;Date: 11-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 17-Mar-1999
C;Accession: $26527; PH1442
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra
J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell
A;Reference number: $26512; MUID:92364546; PMID:1380061
A;Accession: $26527
                                                                                                                                                                                                                                                                                     A; Molecule type: mRNA
A; Restidues: 1-15 < CASS
A; Cross-references: UNIPARC: UPI00001769BB; EMBL: X67978
A; Cross-references: Cytolytic T-lymphocyte, clone Cw3/A8
A; Experimental source: Cytolytic T-lymphocyte, clone Cw3/A8
R; Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pai
J. Exp. Med. 177, 811-820, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Binette, J.P.; Binette, M.B. Scanning Microsc. 8, 233-239, 1994
A;Title: Sequencing of proteins extracted from stones. A;Reference number: A56049; MUID:95215817; PMID:7701298
A;Recession: A56049
                                                                                         A/Experimental source: cytolytic C/Superfamily: immunoglobulin V r C/Keywords: T-cell receptor
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A;Experimental source: urinary tract and gallbladder stones
A;Note: the source is designated as Homo sapiens, however the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     urinary tract stone matrix protein, 40K - unidentified organism (fragment) C;Species: unidentified organism C;Date: 12-Apr-1995 #sequence_revision 12-Apr-1995 #text_change 20-Apr-2000 C;Accession: A56049
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T-cell receptor alpha chain (B28) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PHO775
R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P. Maringari, J. Exp. Med. 174, 1371-132 1001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complexallelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0775
A;Molecule type: mRNA
A;Residues: 1-15 <CAS>
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A;Title: Identification of tiesue-specific
A;Reference number: S43624; MUID:94237150;
A;Accession: S43634
A;Molecule type: protein
A;Residues: 1-15 <FRE>
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                 A;Reference number: PH0746;
A;Accession: PH0784
A;Molecule type: mRNA
                                                            A;Title: T cell receptor genes in a series of class I major histocompatibility complex-allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
                                                                                                                                                                 T-cell receptor alpha chain (F1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0784
                                                                                                                           R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L. J. Exp. Med. 174, 1371-1383, 1991
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C;Keywords: cardiac muscle; heart; membrane-associated complex; mitochondrion; oxidored
                                                                                                                                                                                                                                                                               RESULT
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1-15 <CAS>
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Pred. No. 4.1e+03;
0; Mismatches 2
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. 4.1e+03;
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interphotoreceptor retinoid-binding protein - ha NyAlternate names: interstitial retinol-binding c;Species: Cricetinae gen. sp. (hamster) C;Date: 21-May-1988 #sequence_revision 21-May-19 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417 C;Accession: G24417
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S05699
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A;Residues: 1-15 <CAS>
A;Cross-references: UNIPARC:UPI0000115FAA;
A;Experimental source: T lymphocyte
C;Keywords: T-cell receptor
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A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPARC:UPI000017C76B; EMBL:X60885
A;Experimental source: T lymphocyte
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Roghani, M.; Hossenlopp, P.; Lepage, P.; Balland, A.; Binoux, FEBS Lett. 255, 253-258, 1899
A;Tkle: Isolation from human cerebrospinal fluid of a new insu.
A;Reference number: S05699; MUID:90005986; PMID:2551732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
PH0751
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                                                                                                                                                                                                                              RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-15 <ROG>
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;Species: Mus musculus (house mouse)
;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 05-Nov-1999
;Accession: PH0751
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Matches
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Best Local !
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3; Conserv

 Conservative

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Pred. No. 6.3e+03;
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Pred. No. 6
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                                                                       21-May-1988
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6.3e+03;
                                                                                                                                                                hamster (fragment)
         G.I.;
                                                                                                                               protein
                                                                 #text_change 09-Jul-2004
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      Landers,
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         R.A.; Bridges,
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Lett. 205, 309-312,

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RESULT 15
F29501
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unpublished results, cited by Blombaeck, B., and Blombaeck, M., in Chemota
A;Reference number: A29501
A;Accession: F29501
A;Status: preliminary
A;Molecule type: protein
A;Molecule type: protein
A;Residues: 1-15 - GLO>
A;Cross-references: UNIPROT:Q7M3I8; UNIPARC:UPI00001777D7
A;Cross-references: UNIPROT:Q7M3I8; UNIPARC:UPI00001777D7
                                                                                                                                                                                                                                                                                                                                                                                                                                                             fibrinopeptide A - wombat
C;Specise: Vombatidae gen. sp. (wombat)
C;Date: 21-Nov-1987 #sequence_revision 08-Jun-1990 #text_change 09-Jul-2004
C;Accession: F29501
R;Blombaeck, B.; Blombaeck, M.; Hann, C.
Search completed: June 13, 2006, 02:47:43 Job time : 39 secs
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C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Reference number: PC2214; MUID:94296418; PMID:8024586
A;Accession: PC2215
A;Molecule type: protein
A;Residues: 1-15 <HUN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fibrinogenolytic proteinase A2 (EC 3.4.21.-) - western diamondback rattlesnake (fragment N;Alternate names: alpha-fibrinogenase A2 C;Species: Crotalus atrox (western diamondback rattlesnake) C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
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P14463;
O1-JAN-1990, integrated into UniProtKB/Swiss-Prot.
O1-JAN-1990, sequence version 1.
O7-FEB-2006, entry version 29.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
Pecora; Bovidae; Bov
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Name=FGA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=16091434; DOI=10.1099/jmm.0.45956-0;
Vitorino L., de Sousa R., Bacellar F., Ze-Ze L.;
"Characterization of a tandem repeat polymorphism in Rickettsia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales; Rickettsiaceae; Rickettsieae; Rickettsia; spotted fever group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-SEP-2005, integrated into UniProtKB/TrEMBL.
27-SEP-2005, sequence version 1.
07-FBB-2006, entry version 3.
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Q3ZJF3;
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Submitted (MAR-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=37816;
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  Bovinae;
             Chordata; Craniata; Vertebrata; Euteleostomi; Laurasiatheria; Cetartiodactyla; Ruminantia;
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RESULT 5
Q73362 N
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RA Galagan
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CONTRAIN-74-OR23-1A / FGSC 987;

XX MEDLINE-22598136; PubMed=12712197; DOI=10.1038/nature01554;

XX Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D.,

XX Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Read N.D.,

XX Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Read N.D.,

XX Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Rehman B.,

XX Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Endrizzi M.,

XX Galagan J.E., Calvo S.E., Nielsen C.B., Butler J., Endrizzi M.,

XX Galagan J.E., Calvo S.E., Nielsen D., Kinsey J.A., Braun E.L.,

XX Galagan J.E., Selltremnikoff C.P., Kinsey J.A., Braun E.L.,

XX Werner-Mashburne M., Selltremnikoff C.P., Kinsey J.A., Braun E.L.,

XX Werner-Mashburne M., Selltremnikoff C.P., Kinsey J.A., Braun E.L.,

XX Celter A., Schulte U., Kothe G.O., Jedd G., Mewes H.-W., Staben C.,

XX Zelter A., Schulte U., Kothe G.O., Jedd G., Mewes H.-W., Staben C.,

XX Amacotte E., Greenberg D., Roy A., Foley K., Naylor J.,

XX Mauceli E., Bielke C., Rudd S., Frishman D., Krystofova S.,

XX Macino G., Catcheside D.E.A., Li W., Pratt R. J., Osmani S.A.,

XX Arden O., Plamann M., Seiler S., Dunlap J.C., Radford A., Aramayo R.,

XX Yarden O., Plamann M., Seiler S., Dunlap J.C., Radford A., Aramayo R.,
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Sordariomycetidae; Sordariales; Sordariaceae; Neurospora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Predicted protein. ORFNames=NCU08588.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-DEC-2003, integrated into UniProtKB/TrEMBL 15-DEC-2003, sequence version 1. 07-FEB-2006, entry version 8.
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Arch. Biochem. Biophys. 118:456-467(197).
-!- FUNCTION: Fibrinogen has a double function: yielding monomers polymerize into fibrin and acting as a cofactor in platelet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=5141;
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Doolittle R.F., Schubert D., Schw
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DOMAIN: A long coiled coil structure formed by 3 polypeptide chains connects the central nodule to the C-terminal domains (distal nodules). The long C-terminal ends of the alpha chains fold back, contributing a fourth strand to the coiled coil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PTM: Conversion of fibrinogen to fibrin is triggered by thrombin, which cleaves fibrinopeptides A and B from alpha and beta chains, and thus exposes the N-terminal polymerization sites responsible for the formation of the soft clot.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     aggregation.
SUBUNIT: Heterohexamer; disulfide linked. Contains
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          coagulation; Coiled coil; Direct protein sequencing.
DE 1 15 Fibrinopeptide A.
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MEDLINE=92020861; PubMed=1717978;
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Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
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-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which preliminary data.
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NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                       MEDLINE=80024255; PubMed=226361; van Heuverswyn H., Fiers W.; "Nucleotide sequence of the Hind-C fragment of Simian Virus 40 DNA: comparison of the 5'-untranslated region of wild-type virus and of some deletion mutants."; some deletion mutants.";
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Pan J., Ghosh P.K., Celma M.L., Weissman
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Science 200:494-502(1978).
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STRAIN-NCTC 11427;

MEDLINE=95020593; PubMed=7934882;

Fenoll A., Munoz R., Garcia E., de la Campa A.G.;

Fenoll A. basis of the optochin-sensitive phenotype of pneumococcus:

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Nature 273:113-120(1978).
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Simian virus 40 (SV40).
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                                       J. Virol. 66:6353-6360(1992)
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Distributed under the Creative Commons Attribution-NoDerivs License
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RESULT 10
Q13377 HUMAN
ID Q13377;
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DT 01-NOV-1996, integrated into UniProtKB/TrEMBL.
DT 01-FEB-2005, sequence version 3.
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-!- FUNCTION: Conjugation of reduced glutathione to a wide number of exogenous and endogenous hydrophobic electrophiles.
-!- CATALYTIC ACTIVITY: RX + glutathione = HX + R-S-glutathione.
-!- SUBUNIT: Monomer and homodimer.
-!- SUBUNIT: Monomer and homodimer.
-!- SUBUNIT: MONOMER and homodimer.
-!- SUBUNIT: MONOMER and to the GST superfamily.
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07-FEB-2006, entry version 16.
Glutathione S-transferase (EC 2.5.1.18) (Fragment).
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GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
Interpro; IPR001070; POlyoma_coat2.
Pfam; PF00761; Polyoma_coat2; 1.
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CHAIN 1 >15 Glutathion
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Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
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GO:0004364; F:glutathione transferase activity; NAS
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Pred. No. 3.2e+04;
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Pred. No. 2.
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L; Mismatches
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RESULT 11

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ID Q86WB2;
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D7 01-JUN-2003, s
D7 01-JUN-2003, s
D7 01-JUN-2003, s
D7 01-JUN-2003, s
D7 01-JUN-2006, e
D8 ACetyl-CoA cax
GN Name-ACACA;
OS Homo sapiens (
CC Eukaryota; Meter
OC Mammalia; Euth
OC Momo.

NCBI TaxID=96C
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CR TISSUS=Testis;
RA Travers M.T..
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GO; GO:0003989; F:acetyl-CoA carboxylase activity; IEA
GO; GO:0016874; F:ligase activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUN-2003, integrated into UniProtKB/TrEMBL.
01-JUN-2003, sequence version 1.
07-FEB-2006, entry version 8.
Acetyl-CoA carboxylase-alpha (EC 6.4.1.2) (Fragment).
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MEDLINE=9703253; PubMed=8875892; DOI=10.1007/8003359900246;
Prosser J., Inglis J.D., Condie A., Ma K., Kerr S., Thakrar
Taylor K., Cameron J.M., Cooke H.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                       Similarity
4; Conserv
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15 AA;
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15 AA; 1713 MW;
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1753 MW;
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Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                  25.7%;
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    Mismatches

                                                                                                                  Score 19;
Pred. No.
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Pred. No. 3
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3.9e+04;
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Best Local S
Matches 4
Query Match
Best Local Similarity
                                         EMBL;
                                                                                                                         PubMed=16091434; DOI=10.1099/jmm.0.45956-0; Vitorino L., de Sousa R., Bacellar F., Ze-Z "Characterization of a tandem repeat polymo
                                                                                                                                                                                     Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
Rickettsiaceae; Rickettsieae; Rickettsia; spotted fever group
NCBI_TaxID=786;
                                                                                                                                                                                                                                                    27-SEP-2005, integrated into UniProtKB/TrEMBL.
27-SEP-2005, sequence version 1.
07-FEB-2006, entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amorphophallus hirtus.
Plastid; Chloroplast.
Eukaryota; Viridiplantae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QWTA3 9ARAE PRELIMINARY; PRT; 15 AA. Q9WTA3; 01-NOV-1999, integrated into UniProtKB/TrEMBL.
                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                               3 RICAK
Q3ZJD3_RICAK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chiang T.Y., Peng C.I.; Submitted (MAR-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=atpB;
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07-FEB-2006,
                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                         Rickettsia akari.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chloroplast.
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                                                                                                       Med. Microbiol. 54:833-841(2005).
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                                         AY820045; AAY40584.1; -; Genomic_DNA.
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                               1720 MW;
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66.7%;
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25.7%;
Score 19;
Pred. No.
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Pred. No. 3.9e+04;
1; Mismatches 1
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IEA.
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polymorphism in
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           DB 2;
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RESULT 14
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ID Q3ZJD4 RICRH
AC Q3ZJD4;
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Best Local (
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                                                                                                                                                                                                                 PubMed=16091434; DOI=10.1099/jmm.0.45956-0;
Vitorino L., de Sousa R., Bacellar F., Ze-Ze L.;
"Characterization of a tandem repeat polymorphism in Rickettsia
                                                                                                                                                                                                                                                                                           STRAIN-MC16;
                                                                                                                                                                                                                                                                                                                                                                                         Rickettsia aeschlimannii.
Bacteria; Proteobacteria; Alpi
Rickettsiaceae; Rickettsieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-SEP-2005, integrated into Uni
27-SEP-2005, sequence version 1.
07-FEB-2006, entry version 3.
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NON_TER 15 15
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SEQUENCE
                         EMBL; AY820043; AAY40582.1; -; Genomic_DNA.
NON_TER 15 15
                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=dksA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q3ZJD5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J. Med. Microbiol. 54:833-841(2005).
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Vitorino L., de Sousa R., Bacellar F., Ze-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=3
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Rickettsiaceae; Rickettsieae; Rickettsia; spotted fever group.
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                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Query
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1: /BMC_Celerra_SIDS3/ptodata/2/iaa/5_COMB.pep:*

2: /BMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*

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Result

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ALIGNMENTS

US-09-239-043D-2101 US-09-239-043D-2101 Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Sequence 2101, Application US/09239043D Septicany: Seduce, Alessandro APPLICANY: Schuck, Setchan APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Exchon, Brian D. APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLICANY: Chesnut, Robert APPLI

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CURRENT APPLICATION NUMBER: US/09/255,501
CURRENT FILING DATE: 1999-02-23
NUMBER OF SEQ ID NOS: 211
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 97
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                       ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-255-501-97
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APPLICANT: Harding Fiona
TITLE OF INVENTION: HUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN
TITLE OF INVENTION: HUMANS AND METHODS FOR CONSTRUCTING, IDENTIFYING AND
TITLE OF INVENTION: PRODUCING SUCH PROTEINS
FILE REFERENCE: GC527
CURRENT APPLICATION UMMBER: US/09/255,501
CURRENT FILING DATE: 1999-02-23
NUMBER OF SEQ ID NOS: 211
SOOTWARE: Patentin Ver. 2.1
SEQ ID NO 96
LENGTH: 15
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Best Local S
Matches 6
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Best Local Similarity
Matches 5; Conserv
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APPLICANT: Harding, Fiona
TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN
TITLE OF INVENTION: HUMANS AND METHODS FOR CONSTRUCTING, IDENTIFYING AND
TITLE OF INVENTION: PRODUCING SUCH PROTEINS
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5. 6596525
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Similarity 75.0%;
6; Conservative
                                SKLSMGSG 8
ASLSLGSG
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Pred. No. 70;
0; Mismatches
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                                                                   Score 27; DB 2;
Pred. No. 2.4e+02;
2; Mismatches 1
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Pred. No. 2.4e+02;
2; Mismatches 1
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US-09-060-872A-96
; Sequence 96, Application US/09060872A
; Sequence 96, Application US/09060872A
; Patent No. 683550
; GENERAL INFORMATION:
; APPLICANT: Estell, David
; APPLICANT: Harding, Fiona
; TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN
; TITLE OF INVENTION: HUMANS AND METHODS FOR CONSTRUCTING, IDENTIFYING AND
; TILLE OF INVENTION: PRODUCING SUCH PROTEINS
; FILE REFERENCE: GC527
; CURRENT FILLING DATE: 1998-04-15
; NUMBER OF SEQ ID NOS: 211
; SOFTMARE: Patentin Ver. 2.1
; SEQ ID NO 96
; LENGTH: 15
; TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-060-872A-97
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APPLICANT: Harding, Fiona
APPLICANT: Harding, Fiona
TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN
TITLE OF INVENTION: HUMANS AND METHODS FOR CONSTRUCTING, IDENTIFYING AN
TITLE OF INVENTION: PRODUCING SUCH PROTEINS
FILE REFERENCE: GC527
CURRENT APPLICATION NUMBER: US/09/060,872A
CURRENT FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 211
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
Sequence 96, Application US/09500135C
PATENT NO. 6838269
GENERAL INFORMATION:
APPLICANT: Estell, David A.
APPLICANT: Harding, Fiona A.
APPLICANT: Harding, Fiona A.
TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 97, Application US/09060872A Patent No. 6835550 GENERAL INFORMATION:
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Best Local :
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Matches 5; Conserv
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Pred. No. 2.4e+02;
2; Mismatches 1
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Pred. No. 2.4e+02;
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US-09-768-080-96
Sequence 96, Application US/09768080
Patent No. 6897049
GENERAL INFORMATION:
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Best Local Similarity
"hes 5; Conserv?
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CURRENT APPLICATION NUMBER: US/09/500,135C
CURRENT FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 236
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 97
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PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 236
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 96
LENGTH: 15
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APPLICANT: Estell, David
APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
FILE REFERENCE: GC527C3
CURRENT APPLICATION NUMBER: US/09/768,080
CURRENT APPLICATION NUMBER: US 09/677,822
PRIOR APPLICATION NUMBER: US 09/677,822
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US 09/500,135
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APPLICANT: Harding, Fiona A.

TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/500,135C CURRENT FILING DATE: 2000-02-08
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TYPE: PRT
ORGANISM: Artificial Sequence
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Pred. No. 2.4e+02;
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Pred. No. 2 4e+02;
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APPLICANT: Harding, Fiona
APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
FILE REFERENCE; GC527C3
CURRENT APPLICATION NUMBER: US/09/768,080
CURRENT FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: US 09/677,822
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/500,135
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOCTWARE: Patentin Ver: 2.1
SEQ ID NO 97
LENGTH: 15
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Best Local Similarity
Thes 5; Conserva
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US-09-677-822A-96
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             Sequence 96, Application US/09677822A

Patent No. 6936249

GENERAL INFORMATION:
APPLICANT: Estell, David
APPLICANT: Harding, Fiona
TITLE OF INVENTION: METHODS OF MAKING AN ALTERED IMMUNOGENIC RESPONSE AND
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
FILE REFERENCE: GC527C2
CURRENT FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/09/677,822A
CURRENT FILING DATE: 2000-02-08
PRIOR FILING DATE: 2000-00-08
PRIOR FILING DATE: 2000-00-08
PRIOR FILING DATE: 2000-00-08
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GENERAL INFORMATION:
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Best Local Similarity
Matches 5; Conserv
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PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 96
LENGTH: 15
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TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICATION NUMBER: US 09/060,872
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Pred. No. 2.
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Pred. No. 2.
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2.4e+02;
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US-09-677-822A-97
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Best Local Similarity
Matches 5; Conserva
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                                                                                                                                                                                                                                                                                            US-08-572-951-6
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CURRENT APPLICATION NUMBER: US/09/677,822A
CURRENT FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US 09/500,135
PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOFTWARE: Patentin Ver: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 97
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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NUMBER OF SEQ ID NOS: 240
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 97, Application US/09677822A Patent No. 6936249
                                                                                                                                                                                                                                                       Sequence 6, Application US/08572951 Patent No. 5824790
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Best Local Similarity
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TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Estell, David APPLICANT: Harding, Fio
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ORGANISM: Artificial Sequence
FEATURE:
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APPLICANT:
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                                                                                                       TITLE OF INVENTION: MODIFICATION OF STARCH
TITLE OF INVENTION: SYNTHESIS IN PLANTS
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
STREET: 1100
                                                 ADDRESSEE: CUSHMAN DARBY & CUSHMAN
ADDRESSEE: Intellectual Property Group of
ADDRESSEE: Pillsbury Madison & Sutro LLP
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                                    1100 New York Avenue, N.W
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KNIGHT, MARY E.
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Pred. No. 2.4e+02;
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Pred. No. 2.4e+02;
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Best Local Similarity
Watches 5; Conserv?
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                 COUNTRY: USA
ZIP: 20005-3918
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPAN-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: MODIFICATION TITLE OF INVENTION: SYNTHESIS IN NUMBER OF SEQUENCES: 41 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/263,921
FILING DATE: 21-UW-1994
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Paul N. KOKLIİB
REGISTRATION NUMBER: 16,773
REFERENCE/DOCKET NUMBER: 222957/1.02.15C
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PRIOR APPLICATION UMBER: 08/346,602
PFILING DATE: 29-NOV-1994
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APPLICATION NUMBER: US
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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APPLICATION NUMBER: US/08/572,951 FILING DATE: 15-DEC-1995 CLASSIFICATION: 800
                                                                                                                                                                                                STREET: 1100 New CITY: Washington STATE: DC
                                                                                                                                                                                                                                  ADDRESSEE: CUSHMAN DARBY & CUSHMAN ADDRESSEE: Intellectual Property Group of ADDRESSEE: Pillsbury Madison & Sutro LLP STREET: 1100 New York Avenue, N.W.
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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5824790
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linear
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KNIGHT, MARY E.
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/346,602 FILING DATE: 29-NOV-1994 CLASSIFICATION: 800

08/263,921

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RESULT 15
US-09-239-043D-2113
; Sequence 2113, Application US/09239043D
; Patent No. 6689363
; GENERAL INFORMATION:
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US-09-255-501-98
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INFORMATION FOR SEQ ID NO: 23
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: GC527
CURRENT APPLICATION NUMBER: US/09/255,501
CURRENT FILING DATE: 1999-02-23
NUMBER OF SEQ ID NOS: 211
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 98
LENGTH: 15
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Best Local Similarity
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APPLICANT: Estell, David
APPLICANT: Harding, Fiona
APPLICANT: Harding, Fiona
TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGEN:
TITLE OF INVENTION: HUMANS AND METHODS FOR CONSTRUCTING,
TITLE OF INVENTION: PRODUCING SUCH PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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Patent No. 659652
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               APPLICANT: Sette, Al APPLICANT: Sidney,
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                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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NAME: Paul N. KOKULLB
REGISTRATION NUMBER: 16,773
REFERENCE/DOCKET NUMBER: 222957/1.02.15C
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: 08/2
FILING DATE: 21-JUN-1994
CLASSIFICATION: 800
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Southwood, Scott
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83.3%;
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100.0%; Pred. No. 3.7e+02;
Live 0; Mismatches 0;
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Pred. No. 3.7e+02;
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Query Match
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; ORGANISM: Orthohepadnaviridae hepatitis B virus
US-09-239-043D-2113
Search completed: June 13, 2006, 02:48:37 Job time: 49 secs
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PRIOR FILING DATE: 1997-11-25
PRIOR APPLICATION NUMBER: US 08/820,360
PRIOR FILING DATE: 1997-03-12
PRIOR APPLICATION NUMBER: US 60/013,363
PRIOR APPLICATION NUMBER: US 08/461,603
PRIOR APPLICATION NUMBER: US 08/461,603
PRIOR APPLICATION NUMBER: US 08/461,603
PRIOR APPLICATION NUMBER: US 08/461,603
                                                                                                                                                                                                                                                                                                                            Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 2579
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2113
LENGTH: 15
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APPLICANT:
APPLICANT:
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APPLICANT:
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PRIOR APPLICATION NUMBER: US 08/278,634
PRIOR FILING DATE: 1994-07-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Inducing Cellular Immune Responses to Hepatitis B Virus TITLE OF INVENTION: Using Peptide and Nucleic Acid Compositions FILE REFERENCE: 2060.0060007
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 08/197,484 PRIOR FILING DATE: 1994-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1994-03-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1994-12-01
PRIOR APPLICATION NUMBER: US 08/344,824
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                                                                                       6 KIPMGVGLS 14
                                                                                                                                2 KLSMGSGLN 10
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Epimmune Inc.
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Celis, Esteban
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                                                                                                                                                                              Conservative
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Pred. No. 3.7e+02;
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Result
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Sequence:
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:*
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ALIGNMENTS

Sequence 3, Application US/10612162 Publication No. US20040014145A1 GENERAL INFORMATION: APPLICANT: Dade Behring Marburg GmbH ITTLE OF INVENTION: Carbohydrate deficient transferrin (CDT)-specific TITLE OF INVENTION: annibodies, their preparation and use FILE REFERENCE: 2002/B001 CURRENT APPLICATION NUMBER: US/10/612,162 CURRENT FILING DATE: 2003-07-03 PRIOR APPLICATION NUMBER: 10230550.1 PRIOR APPLICATION NUMBER: 03011334.4 PRIOR FILING DATE: 2003-05-19 NUMBER OF SEQ ID NOS: 4 SOFTWARE: Patentin Ver. 2.1 Sequence 10, Application US/10769514 | Publication No. US20040258695A1 | GENERAL INFORMATION: APPLICANT: Schryvers, Anthony | TITLE OF INVENTION: Transferrin Binding Peptides and Uses Thereof | FILE REFERENCE: 028722-001 | CURRENT APPLICATION NUMBER: US/10/769,514 | CURRENT FILING DATE: 2004-01-30 | PRIOR APPLICATION NUMBER: US 60/444,113 | PRIOR APPLICATION NUMBER: US 60/444,113 | PRIOR APPLICATION NUMBER: US 60/444,113 | PRIOR FILING DATE: 2003-01-31 | NUMBER OF SEQ ID NOS: 86 | SOFTWARE: FastSEQ for Windows Version 4.0 | SEQ ID NO 10 | LENGTH: 15 RESULT 2 US-10-769-514-10 밁 S US-10-612-162-3 SOFTWALL 15 RESULT 1 US-10-612-162-3 Query Match Best Local S Matches LENGTH: 15 TYPE: PRT ORGANISM: Homo 15; 1 SKLSMGSGLNLSEPN 15 Similarity SKLSMGSGLNLSEPN Conservative sapiens 100.0%; Score 74; DB 4; L 100.0%; Pred. No. 3.4e-06; ... Mismatches 0; 0; Length 15; Indels <u>,,</u> Gaps

0;

ORGANISM: Homo sapiens

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APPLICANT: Schryvers, Anthony
TITLE OF INVENTION: Transferrin Binding Peptides and Use
FILE REFERENCE: 028722-001
CURRENT APPLICATION NUMBER: US/10/769,514
CURRENT FILING DATE: 2004-01-30
PRIOR APPLICATION NUMBER: US 60/444,113
PRIOR FILING DATE: 2003-01-31
PRIOR FILING DATE: 2003-01-31
NUMBER OF SEQ ID NOS: 86
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 42
LENGTH: 15
TYPE: PRT
                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/801,990
CURRENT FILING DATE: 2004-03-15
PRIOR APPLICATION NUMBER: US 60/455,037
PRIOR FILING DATE: 2003-03-14
NUMBER OF SEQ ID NOS: 395
SOFTWARE: Patentin version 3.2
SEQ ID NO 180
LENGTH: 15
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                              Sequence 180, Application No. US20 GENERAL INFORMATION:
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Best Local
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APPLICANT: Schulman, Howard
APPLICANT: Becker, Christopher
TITLE OF INVENTION: BIOWARKERS FOR RHEUMATOID ARTHRITIS
FILE REFERENCE: SURR.121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE: OTHER INFORMATION: TbpB-binding peptides
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Pred. No. 0.0091;
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Pred. No. 0.0091;
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APPLICANT: Schulman, Howard

APPLICANT: Schulman, Howard

APPLICANT: Becker, Christopher

FILE REFERENCE: SURR.121

CURRENT APPLICATION UNMBER: US/10/801,990

CURRENT FILING DATE: 2004-03-15

PRIOR APPLICATION NUMBER: US 60/455,037

PRIOR APPLICATION NUMBER: US 60/455,037

PRIOR FILING DATE: 2003-03-14

NUMBER OF SEQ ID NOS: 395

SOFTWARE: Patentin version 3.2

LENGTH: 15
                                                                                                                                                                                                                                   APPLICANT: Grey, Howard M.
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Inducing Cellular Immune Responses to Hepatitis B Virus
TITLE OF INVENTION: Using Peptide and Nucleic Acid Compositions
FILE REFERRICE: 2060.0060007
CURRENT FILING DATE: 2003-09-04
PRIOR APPLICATION NUMBER: US/10/654,601
CURRENT FILING DATE: 1999-01-27
PRIOR APPLICATION NUMBER: US 09/189,702
PRIOR APPLICATION NUMBER: US 09/189,702
PRIOR FILING DATE: 1999-11-25
PRIOR FILING DATE: 1999-11-35
PRIOR APPLICATION NUMBER: US 08/978,291
PRIOR APPLICATION NUMBER: US 08/978,291
PRIOR APPLICATION NUMBER: US 08/978,360
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PRIOR APPLICATION NUMBER: US 08/820,360
PRIOR PILING DATE: 1995-03-12
PRIOR PILING DATE: 1995-06-05
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Publication No.
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Best Local Similarity
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APPLICANT:
                                                                    PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
PRIOR APPLICATION NUMBER: US 08/344,824
PRIOR FILING DATE: 1994-11-23
PRIOR APPLICATION NUMBER: US 08/278,634
PRIOR FILING DATE: 1994-07-21
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
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APPLICANT: Sidney
APPLICANT: South
PRIOR FILING DATE: 1994-03-04
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 2579
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Vitiello, Maria A.
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Pred. No. 0.
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US-10-769-514-16
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                                                                                                                       ; NUMBER OF SEQ ID NOS: 86
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 15
; LENGTH: 15
; TYPE: PRT
; ORGANISM: M. Catarrhalis
US-10-769-514-15
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US-10-769-514-16
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US-10-654-601-2101
                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Schryvers, Anthony
TITLE OF INVENTION: Transferrin Binding Peptides and Uses Thereof
FILE REFERENCE: 028722-001
CURRENT APPLICATION NUMBER: US/10/769,514
CURRENT FILING DATE: 2004-01-30
PRIOR APPLICATION NUMBER: US 60/444,113
PRIOR FILING DATE: 2003-01-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16, Application US/10769514
Publication No. US20040258695A1
GENERAL INFORMATION:
APPLICANT: Schryvers, Anthony
TITLE OF INVENTION: Transferrin Binding Peptides and
FILE REFERENCE: 028722-001
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LENGTH: 15
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Best Local
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Publication No.
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Best Local :
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CURRENT FILING DATE: 2004-01-30
PRIOR APPLICATION NUMBER: US 60/444,113
PRIOR FILING DATE: 2003-01-31
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SOFTWARE: FastSEQ for Windows Version 4.0
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Local Similarity 50.0%;
les 6; Conservative
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MGYGMALSKIN 11
                               MGSGLNLSEPN 15
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                                                               Conservative
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Pred. No. 2.9e
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Pred. No. 6.8e+02;
3; Mismatches 3;
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Pred. No. 1e+03;
                                                             Mismatches
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2.9e+02;
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RESULT 11
US-10-895-064-574
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; Sequence 97, Application US/10924092A
; Publication No. US20050137112A1
; GENERAL INFORMATION:
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TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN HUMANS AND
TITLE OF INVENTION: MUTANT PROTEINS HAVING, IDENTIFYING AND PRODUCING SUCH PROTEIN
FILE REFERENCE: GC527-1
CURRENT APPLICATION NUMBER: US/10/924,092A
CURRENT FILING DATE: 2004-08-23
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 211
SOFTWARE: PATENTIN OS: 211
SOFTWARE: PATENTIN OS: 2.1
SEQ ID NO 96
LENGTH: 15
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US-10-924-092A-96
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Publication No. US20050137112A1
GENERAL INFORMATION:
APPLICANT: Estell, David A.
APPLICANT: Harding, Fiona A.
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Best Local
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Best Local
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CURRENT APPLICATION NUMBER: US/10/924,092A
CURRENT FILING DATE: 2004-08-23
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 211
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APPLICANT: Harding, Fiona A.
TITLE OF INVENTION: MUTANT PROTEINS HAVING LOWER ALLERGENIC RESPONSE IN HUMANS
TITLE OF INVENTION: MUTANT PROTEINS HAVING, IDENTIFYING AND PRODUCING SUCH F
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ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 1e+03;
2; Mismatches
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Pred. No. 1e+03;
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Sequence 574, Application US/10895064 Publication No. US20060018923A1 GENERAL INFORMATION: APPLICANT: PEIRIS, JOSEPH S.M.

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RESULT 13
US-11-020-628-97
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APPLICANT: ESTEEL1, David
APPLICANT: Harding, Fiona
APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCING AND U
FILE REFERENCE: GC527C2
CURRENT APPLICATION NUMBER: US/11/020,628
CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: US 09/500,135
PRIOR APPLICATION NUMBER: US 09/60,872
PRIOR APPLICATION NUMBER: US 09/60,872
PRIOR FILING DATE: 1998-04-15
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US-11-020-628-96
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US-10-895-064-574
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            Sequence 97, Application US/11020628
Publication No. US20050164257A1
GENERAL INFORMATION:
APPLICANT: Estell, David
APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCTIONS OF MAK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 240
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 96
LENGTH: 15
TYPE: PRT
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APPLICANT: LEUNG, FREDERICK C.
TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT INFECTION AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: V0690.0031
CURRENT APPLICATION NUMBER: US/10/895,064
CURRENT APPLICATION NUMBER: US/10/895,064
CURRENT FILLNG DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 2918
SOFTWARE: PREENTIN Version 3.2
SEQ ID NO 574
LENGTH: 15
TYPE: PRT
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Publication No. US20050164257A1
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FILE REFERENCE: GC527C2
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                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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les 5; Conservarium
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ASLSLGSG 12
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                PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND OF MAKING AND USING THE SAME
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Pred. No. 1e+03;
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Sequence 97, Application US/11020602
Publication No. US20060024764A1
GENERAL INFORMATION:
APPLICANT: Estell, David
APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCING AN AI
TITLE OF INVENTION: METHODS OF MAKING AND USERIES OF INVENTION: WETHODS OF MAKING AND USERIES REFERENCE: GCS27C2
CURRENT APPLICATION NUMBER: US/11/020,602
CURRENT APPLICATION NUMBER: US/11/020,602
PRIOR APPLICATION NUMBER: US 09/500,135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/11/020,602
CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: US 09/500,135
PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 96
LENGTH: 15
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CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: US 09/500,135
PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 97
LENGTH: 15
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Publication No. US20060024764A1
GENERAL INFORMATION:
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APPLICANT: Harding, Fiona
TITLE OF INVENTION: PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND
TITLE OF INVENTION: METHODS OF MAKING AND USING THE SAME
FILE REFERENCE: GC527C2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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Pred. No. 1e+C
2; Mismatches
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Pred. No.
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PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: US 09/060,872
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 240
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 97
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-11-020-602-97
Query Match
Best Local Similarity 62.5%; Score 27; DB 6; Length 15;
Best Local Similarity 62.5%; Pred. No. 1e+03;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
DB 1 SKLSMGSG 8
Search completed: June 13, 2006, 03:02:14

Job time: 181 secs
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Result
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Perfect score:
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Maximum DB
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2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

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4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

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6: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

8: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
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ALIGNMENTS

US-11-295-192A-1160

Sequence 1160, Application US/11295192A Publication No. US20060110803A1 GENERAL INFORMATION:

RESULT 1

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; OTHER INFORMATION: Synthetic peptide US-11-295-192A-1160
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US-11-295-192A-1161
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Sequence 1161, Application US/11295192A
Publication No. US20060110803A1
GENERAL INFORMATION:
APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Duljk, Wouter C.
APPLICANT: Oudsmit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof FILE REFERENCE: 0089 WO ON ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
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Best Local S
Matches 5
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TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEQ ID NOS: 1264
SOFTWARE: Patentin version 3.1
SEQ ID NO 1160
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Crucell Holland B.V. APPLICANT: Ter Meulen, Jan H. APPLICANT: Sloctetra, Jelle W. APPLICANT: Puijk, Wouter C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                     7
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Pred. No. 2.1e+02;
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Sequence 1162, Application US/11295192A

Publication No. US20060110803A1

GENERAL INFORMATION:
APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Puijk, Wouter C.
APPLICANT: Puijk, Wouter C.
APPLICANT: Puijk, Wouter C.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT APPLICATION NUMBER: PCT/EP2004/051102
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR APPLICATION SATE: 2004-06-14
Sequence 1163, Application US/11295192A

Publication No. US20060110803A1

GENERAL INFORMATION:

APPLICANT: Crucell Holland B.V.

APPLICANT: Ter Meulen, Jan H.

APPLICANT: Sloctstra, Jelle W.

APPLICANT: Goudsmit, Jan

TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and

FILE REFERENCE: 0089 WO 00 ORD

CURRENT APPLICATION NUMBER: US/11/295,192A

CURRENT FILING DATE: 2005-12-06

PRIOR APPLICATION NUMBER: PCT/EP2004/051102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Synthetic peptide US-11-295-192A-1162
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NUMBER OF SEQ ID NOS: 1264
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1161
LENGTH-17
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 1162
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Best Local (
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TYPE: PRT
ORGANISM: Artificial sequence
PEATURE:
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Local Similarity 55.6%;
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Similarity 55.6%;
5; Conservative
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Pred. No. 2.1e+02;
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Pred. No. 2.1e+02;
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Sequence 1165, Application US/11295192A
PUBLICATION NO. US20060110803A1
GENERAL INFORMATION.
APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Bloctstra, Jelle W.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Anrigenic peptides of SARS col
FILE REFERENCE: 0009 WO 00 ORD
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2006-14
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Best Local Similarity
Matches 5; Conserve
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APPLICANT: Crucell H
APPLICANT: Ter Meul
APPLICANT: Slootstr
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TITLE OF INVENTION: Antigenic peptides of SARS coronavirus
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEQ ID NOS: 1264
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1164
NUMBER OF SEQ ID NOS: 1264
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1165
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SOFTMARE: PatentIn version 3.1
SEQ ID NO 1163
LENGTH: 15
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ORGANISM: Artificial sequence
FEATURE:
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ORGANISM: Artificial sequence
FEATURE:
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Slootstra, Jelle W.
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Pred. No. 2.
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Pred. No. 2.1e+02;
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                                         SOFTWARE: PatentIn version 3.1
SEQ ID NO 232
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                            Sequence 232, Application US/11295192A Publication No. US20060110803A1
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Best Local Similarity 55.6%;
Matches 5; Conservative
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APPLICANT: Ter Meulen, Jan H.
APPLICANT: Sloestra, Jelle W.
APPLICANT: Sloestra, Jelle W.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
                                                                                                                      APPLICANT: Puijk, Wouter C.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
                                                                                                                                                                                                                                                                                         APPLICANT: Crucell Holland B.V. APPLICANT: Ter Meulen, Jan H. APPLICANT: Slootstra, Jelle W.
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                                                                                                     NUMBER OF SEQ ID NOS: 1264
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ORGANISM: Artificial sequence
FEATURE:
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ORGANISM: Artificial sequence
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Local Similarity 55.6%;
hes 5; Conservative
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Slootstra, Jelle W.
Puijk, Wouter C.
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Pred. No. 2.1e+02;
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; FEATURE:
; OTHER INFORMATION: Synthetic peptide
US-11-295-192A-232
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PRIOR FILING DATE: 2005-12-06
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEQ ID NOS: 1264
SOFTWARE: Patentin version 3.1
SEQ ID NO 233
LENGTH-----
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                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 234
                                                                                                                                                                                                                                                                                                                                                                         Sequence 234, Application US/11295192A Publication No. US20060110803A1 GENERAL INFORMATION:
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Best Local :
                                                                                                                               APPLICANT: Ter Meulen, Jan H.
APPLICANT: Sloctstra, Jelle W.
APPLICANT: Sloctstra, Jelle W.
APPLICANT: Puljk, Wouter C.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REPERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEQ ID NOS: 1264
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TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
                                                                                                                                                                                                                                                                                                               APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
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APPLICANT: Tex Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Puijk, Wouter C.
                    ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial sequence FEATURE:
OTHER INFORMATION: Synthetic peptide
                                                           TYPE: PRT
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Pred. No. 4.8e+02;
Pred. No. 4.8e+03;
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Pred. No. 4.8e+02;
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Query Match
Best Local Similarity
Matches 4; Conserv

Conservative

1; Mismatches

3; Indels Length 15

0

Gaps

0

28.4%;

Score 21; DB 7; Pred. No. 4.8e+02;

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APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Buijk, Wouter C.
APPLICANT: Goudemit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEQ ID NOS: 1264
SOFTWARE: Patentin version 3:1
SEQ ID NO 236
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
OTHER INFORMATION: Synthetic peptide
US-11-295-192A-236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Puijk, Wouter C.
APPLICANT: Goudemit, Jaap
TITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 ORD
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: PCT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
NUMBER OF SEO ID NOS: 1264
SOFTWARE: Patentin version 3.1
SEQ ID NO 235
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
ORGANISM: Artificial sequence
FEATURE:
OTHER INFORMATION: Synthetic peptide
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Query Match
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Matches 4; Conserv
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Pred. No. 4.8e+02;
  Score 21; DB 7;
Pred. No. 4.8e+02;
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APPLICANT: Goudsmit, Jaap;
ITILE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof;
FILE REFERENCE: 0089 WO 00 ORD;
CURRENT APPLICATION NUMBER: US/11/295,192A;
CURRENT FILING DATE: 2005-12-06;
PRIOR APPLICATION NUMBER: PCT/EP2004/051102;
PRIOR FILING DATE: 2004-06-14;
NUMBER OF SEQ ID NOS: 1264;
SOFTWARE: Patentin version 3.1
SEQ ID NO 237
LENGTH. 15
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US-11-295-192A-237
                                                                                                          ; OTHER INFORMATION: Synthetic peptide US-11-295-192A-238
                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 1264
SOFTWARE: PatentIn version 3.1
SEQ ID NO 238
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Crucell Holland B.V.
APPLICANT: Ter Meulen, Jan H.
APPLICANT: Slootstra, Jelle W.
APPLICANT: Puijk, Wouter C.
                                  Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 238, Application US/11295192A
Publication No. US20060110803A1
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APPLICANT: Crucell Holland B.V.
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APPLICANT: Sloctetra, Jelle W.
APPLICANT: Puljk, Wouter C.
APPLICANT: Goudsmit, Jaap
IITLE OF INVENTION: Antigenic peptides of SARS coronavirus and used thereof
FILE REFERENCE: 0089 WO 00 CND
CURRENT APPLICATION NUMBER: US/11/295,192A
CURRENT FILING DATE: 2005-12-06
PRIOR APPLICATION NUMBER: ECT/EP2004/051102
PRIOR FILING DATE: 2004-06-14
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TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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                                                                                                                                                               ORGANISM: Artificial sequence
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                                      Conservative
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                                                       Score 21; DB 7;
Pred. No. 4.8e+02;
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Pred. No. 4.8e+02;
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RESULT 15
US-11-295-192A-239
INS-11-295-192A-239
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Result
No.
 Minimum
Maximum
 Title:
Perfect score:
 Database
 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
 Sequence:
 Run on:
 OM protein - protein search, using
 Post-processing:
score
 Pred. No.
 DB DB
 Score
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8
 No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.
length:
 A_Geneseq_8:*
1: geneseqp198
2: geneseqp199
3: geneseqp200
4: geneseqp200
 Query
Match
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6:
9:
10:
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 US-10-612-162A-4
68
 June 13, 2006, 03:50:08 ; Search time 84.1176 Seconds (without alignments)
70.661 Million cell updates/sec
 100.0
 100.0
 2589679 seqs, 457216429 residues
 Gapop 10.0 , Gapext 0.5
 BLOSUM62
 Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
 YEKYLGEEYVKAV 13
 Copyright
 geneseqp1980s:*
geneseqp2000s:*
 geneseqp2001s:*
geneseqp2002s:*
geneseqp2003as:*
 geneseqp2003bs:*
geneseqp2004s:*
geneseqp2005s:*
 geneseqp2006s:*
 Length
 2000000000
GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.
 DB
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ABM83780
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ABM83781
ABM83779
ADT08174
ABM83778
 ADH89394
ADK15882
ADL70756
ADF74798
AAB84371
 ABP72820
ADH89360
ADK15869
ADL70732
ADL24413
 IJ
 ABR82321
 AAU02980
 ABP72836
 ADG46078
 33
 SUMMARIES
 mode1
 2589679
 Adg46078 Human CDT
Abg72836 Human tra
Adh89394 Human tra
Adh89394 Human tra
Adk15882 Human tra
Adf74798 Human tra
Adf74798 Human noc
Aab84371 Amino aci
Aau02980 Angiotens
Abr82321 Human met
Abm83782 Human dia
Abm83781 Human dia
Abm83779 Human dia
Abm83779 Human dia
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Abm83778 Human dia
Abm83778 Human dia
Abm83780 Human tra
Abm83780 Human tra
Adh89360 Human tra
Adh89360 Human tra
Abm83780
Abp72820
Adh89360
Adk15869
Adl70732
Adl24413
Adu17885
 Description
Human
Human
Human
 RESULT 1
ADG46078
WPI; 2004-073743/08.
 Althaus
 05-JUL-2002; 2002DE-01030550.
 (DADE-) DADE
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|-----------|-----------|-----------|-----------|-----------|-----------|----------|-------------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| 68        | 68        | 83        | 83        | 83        | 83        | 83       | 83          | 68        | 83        | 68        | 89        | 83        | 68        | 83        | 83        | 68        | 83        | 83        | 83        | 89        | 83        |
| 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0    | 100.0       | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     | 100.0     |
| 869       | 869       | 869       | 869       | 869       | 869       | 696      | 679         | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       | 679       |
| 7         | σ         | ω         | N         | N         | N         | _        | 10          | 9         | 9         | ø         | ω         | ω         | œ         | ω         | œ         | ω         | ω         | œ         | ω         | ω         | œ         |
| ADD45282  | ABP72819  | AAY50717  | AAW54354  | AAR66492  | AAR12499  | AAP70384 | AEE47501    | ADY57103  | ADY57102  | ADY54149  | ADT55066  | ADU00630  | ADU17468  | ADU17611  | ADT98423  | ADT36538  | ADT99248  | ADT78149  | ADT78912  | ADU07204  | ADT55264  |
| Add45282  | Abp72819  | Aay50717  | Aaw54354  | Aar66492  | Aar12499  | Aap70384 | Aee47501    | Ady57103  | Ady57102  | Ady54149  | Adt55066  | Adu00630  | Adu17468  | Adu17611  | Adt98423  | Adt36538  | Adt99248  | Adt78149  | Adt78912  | Adu07204  | Adt55264  |
| Human Pro | Human tra | Human ser | Serotrans | Human tra | Human tra | Sequence | 1 Human tra | Human ser | Human ser | Amino aci | Human par | Human tra | Human tra | Human tra | Human tra | Partial h | Partial h | Human tra | Human tra | Amino aci | Partial h |

ALIGNMENTS

#### 19-MAY-2003; 2003EP-00011334 07-JAN-2004. EP1378521-A1. antibody; carbohydrate-deficient transferrin; CDT; non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human. Human CDT peptide fragment 25-MAR-2004 ADG46078; ADG46078 standard; peptide; 13 sapiens. (first entry) #4. \$

New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization.

BEHRING MARBURG GMBH

Claim 4; SEQ ID NO 4; 21pp; German.

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences vVARSNGGKEDLIWELL, TTEDSTAKIMNGEADAWSLDGGF, SKLSWGSGLINLSERN and YEKYLGEEYVKAV (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated rransferrin or a fragment than general ton of hybridomas by standard transferrin or a fragment than convertions of without the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the sequence of the s transferrin, or a fragment, of spleen and agment, then generation of hybridomas by standard myeloma cells. Hybridomas are then selected for

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RESULT 2
ABP72836
ID ABP7
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 Query Match
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Matches 13
 production of antibodies that specifically bind CDT from the aqueous phase. The antibodies are useful for diagnosis of alcoholism, CDT, which lacks glycosylation at Asn413 and/or Asn 611, is present only in alcoholics. The antibodies allow direct detection of CDT in solution, eliminating the need for immobilising it on a solid phase (as required when using known antibodies). ADG46075-ADG46094 represent CDT peptide fragments used in the method of the invention.
 The present sequence is a portion of the human transferrin (Tf) C don comprising amino acids 611-659 of the Tf sequence given in ABP72820. invention relates to novel fusion proteins of Tf and a therapeutic protein or peptide. The therapeutic protein or peptide, e.g. an erythropoletin mimetic (see ABP72834), can be fused to the N- or C-terminus of Tf, or inserted into Tf especially between residues Glu-6
 New fusion protein, useful in the diagnosis and treatment of diseases or disorders relating to the respiratory, cardiovascular and digestive systems, comprises a transferrin protein fused to a therapeutic protein.
 Transferrin; neuroprotective; cerebroprotective; vasotropic; antiaffackinsonian; nootropic; anti-HIV; antiafshimatic; antiaffergic; cytostatic; immunosuppressive; antiaffacenic; cardiant; gynaecological; immunostimulant; antianaemic; haemostatic; antiparasitic; antiinflammatory; dermatological; antibacterial; virucide; antiparasitic;
 Example 3;
 WPI; 2003-332916/31.
 Misc-difference
 gastrointestinal;
 ABP72836 standard; peptide; 49
 Prior
 30-AUG-2001;
30-NOV-2001;
 30-AUG-2002; 2002WO-US027637
 13-MAR-2003
 WO2003020746-A1
 fungicide; hepatotropic; antirheumatic; antiarthritic; antigout;
tranquillizer; vulnerary; antidiabetic; nephrotropic; antipyretic;
 Human transferrin
 11-AUG-2003
 Sequence 13 AA;
 Modified-site
 ABP72836,
 (BIOR-) BIOREXIS
 sapiens.
 \vdash
 1 YEKYLGEEYVKAV 13
 Similarity
 YEKYLGEEYVKAV 13
 Page 249;
 Conservative
 2001US-0315745P
2001US-0334059P
 (first entry)
 15.
 Location/Qualifiers
 PHARM CORP
 note= "Asn is N-glycosylated"
 gene therapy; transgenic
 peptide insertion region
 . 16
 100.0%;
 298pp;
 "peptide insertion site"
 o,
 English.
 Score 68;
Pred. No.
 Mismatches
 DB 8; 1
 Length 13;
 Indels
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 Gaps
 domain,
20. The
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ADH89394
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This invention relates to a novel fusion protein which comprises a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region. The invention may be useful for the development of compounds with cytostatic, antibacterial, virucide, antiparasitic, immunosuppressive or antiarthritic activity. In addition, the sequences disclosed may be useful for gene therapy. The fusion protein is useful for preparing a composition for treating a disease or disease symptom in a patient for example septic shock, endotoxic shock, cachexia syndromes associated with bacterial, viral or parasitic infections, neoplasm, autoimmune disease, arthritis or adverse effects associated with treatment for preventing graft rejection. The present
 fusion protein; transferrin protein; glycosylation; antibody variable region; cytostatic; antibacterial; virucide; antiparasitic; immunosuppressive; antiarthritic; gene therapy; septic shock; endotoxic shock; cachexia syndrome; bacterial infectivital infection; parasitic infection; neoplasm; autoimmune disease; arthritis; graft rejection; human.
 and Thr-626 of the Tf N domain. Modified Tf fusion proteins of the invention can be used in the diagnosis, prognosis, prevention and/or treatment of diseases and/or disorders of the endocrine, nervous, immune, respiratory, cardiovascular, reproductive and digestive systems, diseases and/or disorders relating to the blood or to cell proliferation, inflammatory conditions and infectious diseases, or to deliver a therapeutic agent to a cell or across the blood-brain barrier
 New fusion protein comprising a transferrin protein exhibiting reduced glycosylation fused to at least one antibody variable region, useful fi preparing a composition for treating e.g., septic shock, neoplasm or
 30-AUG-2001; 2001US-0315745P
30-NCV-2001; 2001US-0334059
30-AUG-2002; 2002US-00231494
30-AUG-2002; 2002US-0406977P
 US2003226155-A1.
 Homo sapiens
 Human transferrin protein domain SeqID38
 15-APR-2004
 ADH89394;
 ADH89394 standard; protein; 49
 Sequence
 Example 2;
 10-MAR-2003; 2003US-00384060
 (BIOR-) BIOREXIS PHARM CORP
 Local
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 YEKYLGEEYVKAV 13
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 YEKYLGEEYVKAV 49
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 Prior CP,
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 ID NO
 100.0%;
 38;
 82pp; English.
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 Score 68;
Pred. No.
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 Mismatches
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 infection;
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RESULT 4
ADK15882
ID ADK1
XX ADK1
AC ADK1
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KW T-20
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RESULT 5
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 The invention comprises a fusion protein that contains a modified transferrin (Tf) protein fused to a therapeutic protein/peptide (e.g. beta-interferon - IFN, glucagons-like peptide - GLP-1, erythropoietimimetic peptide - EMP1, T-20 and soluble toxin receptor). The fusior protein of the invention is useful for epitope tagging. The present acid sequence represents a fragment of the human Tf protein.
 30-AUG-2001; 2001US-0315745P.
30-NOV-2001; 2001US-0334055P.
30-AUG-2002; 2002US-00231494.
30-AUG-2002; 2002US-0406977P.
 glucagons-like peptide; GLP-1; erythropoietin mimetic peptide; T-20; soluble toxin receptor; epitope tagging; human.
 sequence is used in the
ADL70756 standard; peptide;
 04-MAR-2003;
 Homo sapiens
 fusion protein; transferrin; Tf; beta-interferon; beta-IFN;
 Sequence 49
 Sequence 49
 WPI; 2004-010899/01
 Prior
 US2003221201-A1
 Human transferrin
 06-MAY-2004
 Example
 (BIOR-)
 ADK15882,
 ADK15882 standard; protein; 49
 Local
 9
 fusion protein comprising
 therapeutic protein
 CP,
 37
 l Similarity
 13;
 BIOREXIS
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 Similarity
 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 49
 YEKYLGEEYVKAV
 YEKYLGEEYVKÁV 49
 SEQ ID NO 16; 70pp; English.
 Lai C,
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 that of a domain of the human transferrin protein which was
 Conservative
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 Conservative
 exemplification of the
 2003US-00378094.
 (first
 PHARM CORP.
 (Tf) protein fragment
 Sadeghi
 100.0%;
 100.0%;
 or peptide, useful for
 Ħ
 0
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 Score
Pred.
 Score 68;
Pred. No.
 a modified transferrin
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 Mismatches
 Mismatches
 invention.
 No.
 0.0018;
 #4.
 DB
 DB 8;
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 Length 49;
 epitope tagging.
 Length
 Indels
 Indels
 erythropoietin
 49;
 0
 0
 fusion
 Gaps
 Gaps
 amino
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30-AUG-2002;
10-MAR-2003;
09-JUL-2003;
 Dermatological; Antiinflammatory; Antibacterial; Vasotropic; Nephrotropic; Neuroprotective; Cycostatic; Cerebroprotective; Vulnerary; Antiansemic, Antiahberosclerotic; Antiahberosclerotic; Thrombolytic; Anticoagulant; Gastrointestinal; Respiratory; transferrin;
 03-JUN-2004
 WPI; 2004-239175/22
 28-AUG-2003;
 WO2004020588-A2
 Homo sapiens
 Tf; transferrin fusion protein; Tf fusion protein; human.
 Human transferrin,
 (BIOR-) BIOREXIS PHARM CORP
 Immunosuppressive; Haemostatic; Antiallergic;
 ADL70756;
 Turner AJ,
 2002US-0406977P.
2003US-00384060.
2003US-0485404P.
 2003WO-US026779
 (first entry)
 Τf,
 insertion region SEQ
 Sadeghi
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 Antiasthmatic;
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Novel library containing several fusion proteins each of which comprises first transferrin polypeptide fused to at least one second peptide, useful for screening for transferrin fusion protein having the particula particular

Example 3; SEQ ID NO 27; 243pp; English

CC eczema; inflammatory conditions e.g., inflammation associated with CC infection (e.g., septic shock, sepsis), ischaemia-reperfusion injury, CC nephritis, Crohn's disease, multiple sclerosis, respiratory disorders CC (asthma and allergy), gastrointestinal disorders (inflammatory bowel CC (asthma and allergy), gastric, ovarian, lung, bladder), CNS disorders CC (multiple sclerosis, stroke, traumatic brain injury, neurodegenerative CC disorders such as Parkinson's disease, Alzheimer's disease), etc. The CC fusion protein is also useful as an adjuvant to enhance antibacterial or CC undetermined significance (MGUS), waldenstic immune responses, etc. The CC undetermined significance (MGUS), waldenstrom's disease, plasmacytomas, adult respiratory distress syndrome, for stimulating wound repair, for CC protein is also useful for treating monoclonal gammopathy of CC protein is also useful for treating monoclonal gammopathy of CC protein is also useful for treating thrombosis, myocardial CC ginarction, cancers, thrombocytopenia, sickle cell anaemia, cc atherosclerosis, etc. The present sequence is a DNA sequence for a CC peptide fragment of Tf(611-659) illustrating an insertion site CC (Glu625/Thr626) for producing the Tf fusion proteins. proteins of transferrin (Tf) and therapeutic proteins with increased serum half-life or serum stability. Preferred fusion proteins include those modified so that the Tf moiety exhibits no or reduced glycosylation, iron binding and/or Tf receptor binding. The transferrin fusion proteins are useful for treating, preventing or ameliorating disorders or diseases of endocrine system, nervous system, immune system, respiratory system, cardiovascular system, diseases and/or disorders relating to cell proliferation, and/or diseases or disorders relating to blood. The modified fusion proteins are useful in diagnosis, prognosis, prevention and/or treatment of autoimmune disorders; diseases and disorders of haematopoietic cells (e.g., leukopenia, neutropenia, anaemia and thrombocytopenia); allergic reactions such as allergic asthma, thinitis and anaphlaxis, IgE-mediated allergic reactions such as asthma, thinitis and The present invention relates to a library of modified l fusion

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S
 Matches
 Query Match
Best Local
 20-MAR-2002;
22-MAR-2002;
12-APR-2002;
12-APR-2002;
12-APR-2002;
23-APR-2002;
12-JUN-2002;
14-JUN-2002;
 06-MAR-2002;
08-MAR-2002;
08-MAR-2002;
15-MAR-2002;
19-MAR-2002;
20-MAR-2002;
20-MAR-2002;
 Sequence 49
 human; NOVX; metabolic disorder; diabetes; anorexia; cancer; cardiovascular; infectious; neurodegenerative; immune; haematopoietic disease; dyslipidaemia; anorectic; virucide; nootropic; antiinflammatory; neuroprotective; antilipaemic; anabolic; cardiant; neurogenesis; wound healing; angiogenesis; chromosome mapping; neurogenesis; wound healing; angiogenesis; chromosome mapping;
 preventing, diagnosing or treating NOVX-a osteoarthritis, obesity, atherosclerosis,
 New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, (
 Alsobrook JP, Anderson DW, Boldog FL, Burgess CE, Chaudhuri A; Colman SD, Edinger SR, Ettenberg S, Gangolli EA, Gerlach VL; Gorman L, Guo X, Kekuda R, Li L, Maclachlan T, Malyankar UM; Mezes PS, Miller CE, Millet I, Padigaru M, Patturajan M, Peyman J; Qian X, Rastelli L, Rieger DK, Smithson G, Spytek KA, Stone DJ; Sukumaran S, Vernet CAM, Voss EZ, Zhong M;
 ADF74798 standard;
 26-JUN-2002;
15-AUG-2002;
 Homo sapiens
 26-FEB-2004 (first entry)
 ADF74798
 06-MAR-2003;
 18-SEP-2003.
 WO2003076578-A2
 tissue typing; preventive
 (CURA-)
 2003-697890/66.
DB; ADF74797.
 NOVX protein to treat human pathological conditions (SeqID 12).
 37
 13
1
 1 YEKYLGEEYVKAV 13
 CURAGEN CORP.
 Similarity
 YEKYLGEEYVKAV 49
 2002US-0365985P.
2002US-0366928P.
2002US-0372018P.
2002US-0372022P.
2002US-0374682P.
2002US-0388096P.
2002US-0388096P.
 Conservative
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 2002US-0412957P.
2002US-0420382P.
2003US-00420382.
 2002US-0365884P.
2002US-0365884P.
2002US-0365984P.
 2002WO-US007288.
2002US-0365034P.
 2003WO-US006794.
 2002US-0361974P.
2002US-00093463.
 protein;
 100.0%;
 100.0%;
 medicine; pharmacogenomic.
 137 AA
 0
 Score 68;
Pred. No.
 Mismatches
 DB 8;
 cancer,
 <u>,,</u>
 Length 49;
 d disorders, e.g.
Parkinson's disease,
 Indels
 0
 Gaps
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 This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical CC thereof, which have properties related to the stimulation of biochemical CC Specifically, it refers to the use of biologically active fragments for CC diagnostic and prognostic assays and furthermore in the treatment of CC diverse pathological conditions. The present invention describes novel human and murine NOVX proteins, as well as methods to modulate their CC expression using antisense oligos, ribozymes and peptide nucleic acids. CC manufacture of a medicament for treating metabolic disorders, diabettes, anorexia, cancer, cardiovascular, infectious, neurodegenerative, immune CC and haematopoietic diseases as well as various dyslipidaemias.

CC virucide, nootropic, antiinflammatory, neuroprotective, antilipaemic, CC virucide, nootropic, antiinflammatory, neuroprotective, antilipaemic, condentify small molecules that modulate or inhibit, for example, conditions and acadisant. Furthermore, they are useful in screening assays to identify small molecules that modulate or inhibit, for example, conserving and anylogenesis. The nucleic acids are also used as in chromosome mapping, tissue typing, preventive medicine and convention.
 Matches
 Query Match
 hyperproliferative disorder; neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebral ischemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; coular disorder; corneal infection; wound healing;
 Claim 11; Page 286-287; 290pp; English.
 Isolated nucleic acid molecule encoding a human tranferrin protein used in preventing, treating or ameliorating a medical condition.
 WO200146254-A1.
 Homo sapiens.
 Amino acid sequence of a human transferrin.
 22-AUG-2001
 AAB84371 standard; protein; 328
 Sequence 137 AA;
 Claim
 Ruben SM,
 21-DEC-2000; 2000WO-US034769.
 28-JUN-2001.
 Human; transferrin; autoimmune disease; rheumatoid arthritis;
 (HUMA-) HUMAN GENOME SCI INC
 Local
 2001-381910/40.
 <u>سر</u>
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 1 Similarity
13; Conserv
 1 YEKYLGEEYVKAV 13
 AAH25190
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 YEKYLGEEYVKAV 117
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 99US-0171595P.
 NO 12; 282pp; English
 100.0%;
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 Score 68;
Pred. No.
 Mismatches
 0.0051;
 DB 7;
 Length 137;
 Indels
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 Gaps
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RESULT 8
ANUO2980
ID ANUO2980
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AC AAUO
AC AAUO
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 The present sequence represents human transferrin polypeptide.

Transferrin polypeptides and polynuclectides are used to prevent, treat
or ameliorate a medical condition in e.g. humans, mice, rabbits, goats,
horses, cats, dogs, chickens or sheep. Disorders which are diagnosed or
treated include autoimmune diseases e.g. rheumatoid arthritis,
hyperproliferative disorders e.g. copplasms of the breast or liver,
cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders
e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g.
Alzheimer's disease, infections caused by bacteria, viruses and fungi and
coular disorders e.g. corneal infection. The polypeptides can also be
used to aid wound healing and epithelial cell proliferation, to prevent
skin aging due to sumburn, to maintain organs before transplantation, for
supporting cell culture of primary tissues, to regenerate tissues and in
chemotaxis. The polypeptides can also be used as a food additive or
preservative to increase or decrease storage capabilities
 Matches
 Query Match
Best Local
 Angiotensin converting enzyme splice variant; ACEV; interleukin 6; granulocyte colony stimulating factor receptor; glucagon; hypertrophy; platelet-derived endothelial cell growth factor; cardiovascular disease; cellular tumour antigen P53; cyclin-dependent kinase inhibitor 10; vasoactive intestinal polypeptide receptor 2; arteriosclerosis; cancer; myocardial infarction; coronary arterial thrombosis; renal disease; miorardiosis; immune disease; immune disease; adiabetic nephropathy; muscular disease; immune disease; endothelial abnormality; nonarcoidotic pulmonary granulomatous disease; endothelial abnormality;
 Novel alternative splicing variants e.g. variant of angiotensin converting enzyme (ACEV), useful in identifying candidate compounds capable of binding to the variant and to detect anti-variant antibo
 17-NOV-1999;
10-DEC-1999;
 Claim 4; Fig
 WO200136632-A2
 Angiotensin converting enzyme (ACEV) splice variant protein #80
 AAU02980 standard; protein; 539
 Sequence 328
 Levine
 17-NOV-2000; 2000WO-IL000766
 25-MAY-2001.
 vascular disorder; asbestosis.
 (COMP-)
 2001-336004/35.
 296 YEKYLGEEYVKAV 308
 13; Conserv
 1 YEKYLGEEYVKAV 13
 COMPUGEN LTD
 AAS06080.
 David
 Conservative
 80; 519pp; English
 8
 (first entry)
 99IL-00132978.
99IL-00133455.
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 100.0%;
 Azar
 0
 Score 68;
Pred. No.
 Khosravi R,
 Mismatches
 , DB 4;
. 0.013;
 Bernstein J;
 0
 Length 328;
 Indels
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 antibodies.
 Gaps
 0
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RESULT 9
ABR82
ID ABR88
XX ABR82
AC ABR8
XX ABP82
DT 06-)
XX MEDP1
KW MEDP1
KW Anti
KW Anti
KW Anti
KW Anti
KW Anti
KW Anti
KW Anti
KW Anti
KW Anti
EMB AC ABR8
XX MEDP1
PR 18--
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 CC (ACEV) polypeptide. The polypeptides of the invention include variants of CC granulocyte colony stimulating factor receptor, glucagon, interleukin 6, CC platelet-derived endothelial cell growth factor, cyclin-dependent kinase CC inhibitor 1C, cellular tumour antigen P53, and vasoactive intestinal CC polypeptide receptor 2. The polypeptides and their associated nucleic CC candidate compounds capable of binding the molecules. The sequences of CC candidate compounds capable of binding the molecules. The sequences of the invention can be used in the treatment and diagnosis of various CC disorders including cardiovascular diseases such as arteriosclerosis, CC myocardial infarction and coronary arterial thrombosis, renal diseases CC such as diabetic nephropathy, muscular diseases such as hypertrophy, CC immune disorders such as immune complex nephritis, multiple sclerosis, CC cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous diseases such as abestosis and vascular pathologies involving an endothelial CC abnormality such as deep vein thrombosis
 Matches
 Query Match
 MEPR; metalloprotein; anti-HIV; antiallergic; antiinflammatory; human; antianemic; antiparkinsonian; nootropic; anticonvulsant; cytostatic; antiarteriosclerotic; antiasthmatic; immunosuppressive; antitityroid; hepatotropic; dermatological; antidiabetic; nephrotropic; antigout; thyromimetic; neuroprotective; osteopathic; antiarthritic; uropathic; antiparasitic; antihelmintic; antipsoriatic; ophthalmological; virucide; antiheumatic; haemostatic; antibacterial; protozoacide; fungicide;
 Kable AE,
Emerling
 14-JAN-2002; 2002US-0348769P.
18-JAN-2002; 2002US-0350701P.
19-MAR-2002; 2002US-0366059P.
10-MAY-2002; 2002US-0379907P.
 Human metalloprotein (MEPR) polypeptide (Id-7509328CD1).
 Sequence
 N-PSDB; ACF35812.
 WO2003060089-A2.
 ABR82321 standard; protein; 575
 The sequence represents an angiotensin converting enzyme splice variant (ACEV) polypeptide. The polypeptides of the invention include variants (
 WPI; 2003-598523/56.
 14-JAN-2003; 2003WO-US001485
 24-JUL-2003.
 (INCY-) INCYTE GENOMICS
 Local
 l similarity
13; Conserv
 BM,
 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 519
 539 AA;
 Griffin JA,
M, Chien D,
 Conservative
 (first entry)
 100.0%;
 Gorvad
Jin P,
 INC.
 0,
 Score 68; DB 4;
Pred. No. 0.021;
 AE, Becha SD, I
Chawla NK, Yue
 Mismatches
 4.
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 Richardson
e H, Khare
 Length 539;
 Indels
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 Marquis JP;
 Gaps
 0
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New human metalloproteins and polynucleotides, useful for diagnosing treating or preventing autoimmune or inflammatory disorders (e.g. All allergy or anemia), multiple sclerosis, osteoarthritis, cancer or

Claim

1; Page 146-148; 153pp; English

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RESULT 10
ABM83782
ID ABM83
XX ABM83
XX ABM83
XX ABM83
XX ABM83
XX Gene
XX Gene
XX Gene
XX WO20C
XX 25-MJ
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XX L12-SI
XX L12-SI
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 The invention relates to novel human metalloproteins (MEPR) and encoding CC polynucleotides. The human MEPR polypeptides, polynucleotides and cc modulators are useful for diagnosing, treating or preventing disorders associated with aberrant expression of MEPR, particularly cell CC proliferative disorders (e.g. arteriosclerosis, atherosclerosis, pepatitis, paroxysmal nocturnal hemoglobinuria, polycythemia CC cirrhosis, hepatitis, paroxysmal nocturnal hemoglobinuria, polycythemia CC vera, psoriasis, primary thromobocytopenia or cancer), developmental CC disorders (e.g. renal tubular acidosis, anemia or mental retardation), CC neurological disorders (e.g. Alzheimer's disease, Parkinson's disease or epilepsy), autoimmune/inflammatory disorders (e.g. AIDS, allergies, CC asthma, autoimmune/inflammatory disorders (e.g. AIDS, allergies, CC diabetes mellitus, glomerulonephritis, Goodpasture's syndrome, GC disbetes mellitus, glomerulonephritis, Goodpasture's syndrome, CC diabetes mellitus, osteoporosis, pancreatitis, Reiter's Syndrome, rheumatoid arthritis, osteoporosis, pancreatitis, or viral, CC syndrome, rheumatoid arthritis, Sjogren's syndrome, uveitis, or viral, CC bacterial, fungal, parasitic, protozoan or helminthic infections. The CC polymacleotides encoding MEPR are useful for creating transgenic animals CC to model human disease. Sequences ABR82316-323 represent the human MEPR
 Query Match
Best Local S
Matches 13
 WPI; 2004-329368/30.
N-PSDB; ACN42434.
 Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES;
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietzei
 gene therapy; human diagnostic and therapeutic polynucleotide; dithp
 New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human
 WO2004023973-A2
 Human diagnostic and therapeutic pprotein SEQ ID NO:4031.
 18-NOV-2004
 Patury S,
 12-SEP-2002; 2002US-0410259P
12-SEP-2002; 2002US-0410260P
 Homo sapiens
 ABM83782;
 Sequence 575
 polypeptides of the invention
 (INCY-) INCYTE CORP.
 12-SEP-2003; 2003WO-US028227.
 25-MAR-2004
 ABM83782 standard; protein;
 543
 Kwong M, E
S, Shi X,
 13;
 1 YEKYLGEEYVKAV 13
 Similarity
 YEKYLGEEYVKAV 555
 Conservative
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 Policky JL,
Suarez CJ;
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 100.0%;
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 Score 68; DE
Pred. No. 0.0
0; Mismatches
 DB 6;
 0,
 Length 575;
 Indels
 0
 Gietzen D;
 Gaps
 0
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RESULT 11
ABM83781
ID ABM83781
AC ABM83
XX ABM83
XX ABM83
XX ABM83
XX IB-NO
DE Human
XX Gene
XX Gene
XX Homo
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 The invention relates to novel diagnostic and therapeutic polynucleotides celected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human conditions diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders, associated autoimmune/inflammatory disorders, gastrointestinal disorder, endocrine confidence of the conditions caused by virus, bacteria, fungi or parasite. The dithp conditions caused by virus, bacteria, fungi or parasite. The dithp conditions caused by virus, bacteria, fungi or parasite. The dithp conditions caused by virus, bacteria, fungi or parasite. The dithp conditions caused by virus, bacteria, fungi or parasite. The dithp conditions caused by virus, bacteria, fungi or parasite. The dithing conditions also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this pattent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
 Matches
 Best
 Query Match
 scnmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV; Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH; Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES; Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gletzer Patury S, Shi X, Suarez CJ;
New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
 WPI; 2004-329368/30.
N-PSDB; ACN42433.
 12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
 gene therapy; human diagnostic and therapeutic polynucleotide; dithp
 Human diagnostic and therapeutic pprotein SEQ ID NO:4030
 18-NOV-2004
 ABM83781;
 ABM83781 standard; protein; 627
 Sequence 609 AA;
 in gene mapping
 molecules, e.g. autoimmune or inflammatory disorders, in gene therapy
 12-SEP-2003; 2003WO-US028227
 25-MAR-2004.
 WO2004023973-A2
 Homo sapiens
 (INCY-) INCYTE CORP.
 Local
 577
 l Similarity
13; Conserv
 1 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 589
 Page; 190pp; English.
 100.0%; ilarity 100.0%; Conservative 0,
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 entry)
 0
 Score 68;
Pred. No.
 Mismatches
 0.024;
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 Length
 Indels
 609;
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gene therapy or

Claim 27; Page; 190pp; English

in gene mapping

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RESULT 12
ABM83779
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 CC The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine CC disorder, neurological disorders, gastrointestinal disorder, endocrine CC disorder, neurological disorders, gastrointestinal disorder, endocrine CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the CC invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly CC from WIPO at www.wipo.int/pct/en/sequences/listing.htm
 Query Match
Best Local S
Matches 13
 Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietz
Patury S, Shi X, Suarez CJ;
 New diagnostic and therapeutic polynucleotides and polypeptides, use in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therap
 ä
 12-SEP-2002; 2002US-0410259P
12-SEP-2002; 2002US-0410260P
 25-MAR-2004.
 WO2004023973-A2
 Homo sapiens
 gene therapy; human diagnostic and therapeutic polynucleotide; dithp
 Human diagnostic
 18-NOV-2004
 ABM83779;
 ABM83779 standard;
 Sequence 627 AA;
 12-SEP-2003; 2003WO-US028227
gene
 2004-329368/30
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 ACN42431.
 INCYTE CORP.
 Similarity
 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 607
 Conservative
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 protein;
 100.0%;
 therapeutic pprotein SEQ ID NO:4028
 628
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 Score 68; DB 8; Pred. No. 0.024;
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 Mismatches
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 Length 627;
 Indels
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RESULT 1:
ADTO8174
ID ADTO8174
XX ADTO
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XX MBP,
XX MBP,
XX ADTO
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 The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be CC used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine autoimmune/inflammatory disorders, developmental disorders, or CC disorder, neurological disorders, gastrointestinal disorders, or CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the convention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly cc from WIPO at www.wipo.int/pct/en/sequences/listing.htm
 Query Match
Best Local S
Matches 13
 New human metal binding proteins (MBP), useful for diagnosing, treating and preventing diseases or conditions associated with the aberrant MBP expression e.g. cancer, AIDS, epilepsy, or infections.
 N-PSDB;
 MBP; metal binding protein; cytostatic; antiarteriosclerotic; anti-HIV; anticonvulsant; nootropic; neuroprotective; cerebroprotective; anti-HIV; antiallergic; antiinflammatory; thyromimetic; transgenic; human.
 Human metal binding
 24-MAR-2003; 2003US-0457538P
 17-MAR-2004; 2004WO-US008328.
 WO2004085612-A2.
 Homo sapiens
 30-DEC-2004
 ADT08174;
 ADT08174
 Sequence
 Claim 27; Page; 190pp; English
 2004-710307/69.
DB; ADT08177.
 596 YEKYLGEEYVKAV 608
 ×
 l Similarity
13; Conserv
 standard; protein;
 YEKYLGEEYVKAV 13
 628 AA;
 Jackson AA,
 100.0%; ilarity 100.0%; Conservative 0
 (first entry)
 protein (MBP)
 Hafalia
 635 AA
 <u>.</u>
 Score 68;
Pred. No.
 ĀJА,
 Mismatches
 (ID:
 Mason
 7512697CD1).
 0.024;
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 8
 0
 Length 628;
 Indels
 <u>.</u>
 Gaps
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The invention relates to novel human metal binding proteins (MBP) and encoding polynucleotides. The MBPs can be expressed by standard recombinant methodology. The polypeptides and polynucleotides are useful in diagnosing, treating and preventing diseases or conditions associated with the decreased expression or overexpression of MBP, such as cell

Claim

SEQ ID

NO 3; 161pp; English

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RESULT 14
ABM033778
ID ABM03
XX
AC ABM03
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AC ABM03
XX
DT 18-N(
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GEN
COS Homc
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FO
 CC proliferative (e.g. cancer, atherosclerosis), neurological (e.g. AIDS, CC epilepsy, Huntington's disease, stroke), immune/inflammatory (e.g. AIDS, CC allergies) and developmental (e.g. Hypothyroidism, Cushing's syndrome) CC disorders, or infections. These are also useful in assessing the effects CC of exogenous compounds on the expression of nucleic acid and amino acid CC sequences of MBP. The MBP or its fragments are useful in screening CC compounds for effectiveness as agonist or antagonist of the polypeptides, or in altering the expression of the target polynucleotide and compounds CC that specifically bind to or modulate the activity of the polypeptide. An CC microarray comprising different nucleotide molecules affixed at distinct CC physical locations on a solid substrate, where at least one nucleotide completes affirst oligonucleotide or polynucleotide sequence CC specifically hybridizable with at least 30 contiguous nucleotides of the carget polynucleotide is useful in monitoring or measuring proteinfunction interactions, drug-target interactions, and gene expression
 Query Match
Best Local S
Matches 13
 Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietz;
Patury S, Shi X, Suarez CJ;
 New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or
 WPI; 2004-329368/30.
N-PSDB; ACN42430.
 12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P.
 gene therapy; human diagnostic and therapeutic polynucleotide; dithp
 Claim 27; Page; 190pp; English.
 25-MAR-2004
 WO2004023973-A2
 18-NOV-2004 (first entry)
 Sequence 635 AA;
 profiles. The present sequence represents a human MBP.
 in gene mapping.
 12-SEP-2003; 2003WO-US028227
 Human diagnostic and therapeutic pprotein SEQ ID NO:4027
 ABM83778
 (INCY-) INCYTE CORP.
 ABM83778 standard; protein; 643 AA.
 603
 100.0%;
al Similarity 100.0%;
13; Conservative 0
 1 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 615
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 Score 68;
Pred. No.
 Mismatches
 DB 8;
 0; Indels
 Length 635,
 in gene therapy or
 Kirton ES;
JL, Gietzen D;
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 Gaps
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The invention relates to novel diagnostic and therapeutic polynucleotides

The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A

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ABM63783
ID ABM6
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 RESULT 15
 CC selected from one of the 2722 sequences defined in the specification. A CC polynuclootide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be CC used to diagnose a particular condition, disease or disorder associated CC autoimmune/inflammatory disorder; developmental disorder, endocrine CC disorder, neurological disorders, developmental disorders, or CC infections caused by virus, bacteria, fungi or parastre. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide CC polymorphisms, as molecular weight markers, and for somatic or germline CC invention. Note: The sequence represents a dithp protein of the CC invention. Note: The sequence data for this patent is not represented in CC the printed specification, but was obtained in electronic format directly CC from WIPO at www.wipo.int/pct/en/sequences/listing.htm
 Query Match
Best Local
 Matches
 Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Mooney EM, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES;
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietzer
 New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or in gene mapping.
 Claim 27; Page; 190pp; English
 WPI; 2004-329368/30.
N-PSDB; ACN42435.
 Patury S,
 Schmidt JP,
 12-SEP-2002; 2002US-0410259P
12-SEP-2002; 2002US-0410260P
 12-SEP-2003; 2003WO-US028227.
 25-MAR-2004
 WO2004023973-A2
 gene therapy; human diagnostic and therapeutic polynucleotide; dithp.
 Human diagnostic and therapeutic pprotein SEQ ID NO:4032
 18-NOV-2004
 ABM83783;
 ABM83783 standard;
 Sequence 643 AA;
 Homo sapiens
 (INCY-) INCYTE CORP.
 13;
 Kwong M, I
S, Shi X,
 1 YEKYLGEEYVKAV 13
 Similarity
 YEKYLGEEYVKAV 623
 Conservative
 (first entry)
 Wright RJ,
 Policky JL,
;, Suarez CJ;
 jht RJ, Bruns CM, Marjanovic MM, Shen F;
Suchorolski MT, Altus CM, Pitts SJ, Elder L
Suchorolski MT, Altus CM, Pitts SJ, Reddy TF;
 protein;
 100.0%; Score 68; DB 8; 100.0%; Pred. No. 0.025;
 0
 Mismatches
 0
 Length 643;
 Indels
 Shen c, c.T. Elder LV;
 0
 Gietzen D;
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CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorders.
CC autoimmune/inflammatory disorder, developmental disorder, endocrine CC disorder, neurological disorder, developmental disorders, endocrine CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide coplymorphisms, as molecular weight markers, and for somatic or germline CC invention. Note: The sequence atta for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm

CC from WIPO at www.wipo.int/pct/en/sequences/listing.htm

Query Match
Best Local Similarity 100.0%; Score 68; DB 8; Length 645;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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| 666660000000000000000000000000000000000                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | ngth Di             | * * * * * * or ec                          | )000000<br>Match<br>Match<br>first                          | qs, 9<br>sfyin                 | ດູ                   | -162A<br>EYVKA                | 2006,                                                                              | rch, u       |                                             |
| 1 TFHUP<br>1 TFRUP<br>2 A45543<br>2 JC2323<br>3 49166<br>4 A360166<br>1 TFBOL<br>1 TFHUL<br>1 S01384<br>2 S33761<br>1 A28438<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E75153<br>2 E7515 | ₽ £                 | resul                                      | 000<br>0%<br>100%<br>45 gummari                             | 216763 x<br>chosen             | Gapext 0.5           | -4<br>V 13                    | 04:18:14                                                                           | using sw m   | nCore<br>1993                               |
| transfer transfer lactofer lactofer cribonic lactotra hemiferr lactotra hemiferr lactotra transfer lactotra transfer lactofer UDPJuco glyceral class I glycogen hypothet hypothet bypothet bypothet bypothet proton/s asparagi asparagi asparagi                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | MARIES  Description | predicted by char grown of the rest        | ries                                                        | residues<br>parameters: 283416 |                      |                               | ; Search time 13.1912 Seconds (without alignments) 94.822 Million cell updates/sec | model        | version 5.1.9<br>- 2006 Biocceleration Ltd. |

| 45                 | 44                 | 43                 | 42                | 41                 | 40                 | 39                 | 38                 | 37                 | 36                 | υ<br>5             | 34                 | 33                | 32                 | 31                 | 30                 |
|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|--------------------|
| 39                 | 39                 | 39                 | 39                | 39                 | 39                 | 39                 | 39                 | 39                 | 39                 | 39                 | 39.5               | 40                | 40                 | 40                 | 40                 |
| 57.4               | 57.4               | 57.4               | 57.4              | 57.4               | 57.4               | 57.4               | 57.4               | 57.4               | 57.4               | 57.4               | 58.1               | 58.8              | 58.8               | 58.8               | 58.8               |
| 883                | 731                | 728                | 661               | 607                | 607                | 605                | 600                | 407                | 334                | 311                | 328                | 738               | 635                | 518                | 467                |
| 12                 | N                  | 2                  | N                 | N                  | N                  | N                  | 2                  | N                  | _                  | N                  | ٠,                 | -                 | N                  | _                  | N                  |
| T06848             | D71332             | \$59964            | A69252            | G97928             | B95060             | 139837             | A49230             | E70309             | DEQYG              | A28446             | B64478             | TFHUM             | A42988             | AHMIYX             | 896829             |
| phycobilisome link | probable DNA topoi | procollagen-lysine | 3-hydroxyacyl-CoA | hypothetical prote | dnaK protein [impo | dnaK-type molecula | dnaK-type molecula | hypothetical prote | glyceraldehyde-3-p | transferrin - mous | hypothetical prote | melanotransferrin | dnaK-type molecula | homoserine O-acety | probable sugar tra |

ALIGNMENTS

# A;Cross-references: UNIPROT:P02787; UNIPARC:UPI000002F089; EMBL:M12530; NID:g339452; PID:A;Note: the authors translated the codon CAA for residue 203 as Glu R;MacGillivray, R.T.A.; Mendez, E.; Shewale, J.G.; Sinha, S.K.; Lineback-Zins, J.; Brew, J. Biol. Chem. 258, 3543-3553, 1983 A;Title: The primary structure of human serum transferrin. The structures of seven cyanoc A;Reference number: A92417; MUID:83160878; PMID:6833213 transferrin precursor [validated] - human N;Alternate names: siderophilin C;Species: Homo sapiens (man) C;Date: 15-Oct-1982 #sequence revision 30-Sep-1993 #text\_change 09-Jul-2004 C;Accession: A20981; A92417; Ā94044; A29090; A32739; I51999; I63133; I54011; I68160; A03 R;Yang, F.; Lum, J.B.; McGill, J.R.; Moore, C.M.; Naylor, S.L.; van Bragt, P.H.; Baldwin Proc. Natl. Acad. Sci. U.S.A. 81, 2752-2756, 1984 A;Title: Human transferrin: cDNA characterization and chromosomal localization. A;Reference number: A20981; MUID:84194084; PMID:6585826 A;Contents: variant C A;Molecule type: protein A;Residues: 20-263,'E',265-328,'N',330-379,'SD',382-435,'D',437-557,'T',559-560,'P',562-A;Cross-references: UNIPARC:UPI0000174483 A;Molecule type: mRNA A;Residues: 1-698 <YAN> A; Accession: A20981

A; Note: the sequence shown is the predominant electrophoretic genetic variant (C or TfC)

R;Park, I.; Schaeffer, E.; Sidoli, A.; Baralle, F.E.; Cohen, G.N.; Zakin, M.M. Proc. Natl. Acad. Sci. U.S.A. 82, 3149-3153, 1985
A;Title: Organization of the human transferrin gene: direct evidence that it originated A;Reference number: A94044, MUID:85216459; PMID:3858812
A;Accession: A94044

A;Molecule type: DNA
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A;Residues: 73-263,'E',265-328,'N',330-562 <PAR>
A;Cross-references: UNIPARC:UPI0000174484; EMBL.M11361
A;Adrian, G.S.; Korinek, B.W.; Bowman, B.H.; Yang, F.
Gene 49, 167-175, 1986
A;Title: The human transferrin gene: 5' region contains conserved sequences which match
A;Reference number: A29090; MUID:87192006; PMID:3106157
A;Accession: A29090

A;Molecule type: DNA
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A;Cross-references: UNIPARC:UPI000016B0AC; UNIPARC:UPI000016B0AD; EMBL:M15673
A;Cross-reference, M.; Park, I.; Besmond, C.; Maessen, G.; Trepat, J.S.; Zakin, M.M.; Kalliothem. Biophys. Res. Commun. 119, 273-281, 1984
B;Chem. Biophys. Res. Commun. 119, 273-281, 1984
A;Title: Molecular cloning and sequence analysis of cDNA for human transferrin.
A;Reference number: A32739; MUID:84153910; PMID:6322780
A;Accession: A32739
A;Accession: A32739

A;Molecule type: mRNA A;Residues: 422-690,'G',692-698 <UZA> A;Residues: 422-690,'G',692-698 <UZA> A;Cross-references: UNIPARC:UPI000016B0AB; EMBL:M12525; NID:g339468; PIDN:AAA61142.1; PII R;MacGillivray, R.T.A.; Mendez, E.; Sinha, S.K.; Sutton, M.R.; Lineback-Zins, J.; Brew, I

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transferrin precursor - rabbit
C;Species: Oryctolegue cuniculus (domestic rabbit)
C;Date: 30-Sep-193 #sequence revision 30-Sep-193 #text change 22-Jun-1999
C;Accession: S16246; A61339; C61573; S00335; S02694; A26504; S14853
R;Banfield, D.K.; Chow, B.K.C.; Funk, W.D.; Robertson, K.A.; Umelas, T.M.; W.Biochim. Biophys. Acta 1089, 262-265, 1991
A;Title: The nucleotide sequence of rabbit liver transferrin cDNA.
A;Reference number: S16246; MUID:91274362; PMID:2054387
 A;Cross-references: GDB:120432; OMIM:190000

A;Map position: 3921-3921

A;Introns: 15/1; 72/3; 119/1; 168/1; 212/2; 231/1; 290/3; 350/1; 401/3; 433/1; 444/1; 49

A;Introns: 15/1; 72/3; 119/1; 168/1; 212/2; 231/1; 290/3; 350/1; 401/3; 433/1; 444/1; 49

C;Function:
A;Description: binds iron for delivery into cells

C;Superfamily: transferrin; transferrin repeat homology
C;Superfamily: transferrin; glycoprotein; iron transport; metal binding; plasma
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A;Residues: 636-696 <RE2>
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A;Cross-references: UNIPARC:UPI000016B13C; GB:M26641; NID:g339988; PIDN:AAA61233.1; PID:
R;Schaeffer, E.; Lucero, M.A.; Jeltsch, J.M.; Py, M.C.; Levin, M.J.; Chambon, P.; Cohen,
Gene 56, 109-116, 1987
A;Title: Complete structure of the human transferrin gene. Comparison with analogous chi
A;Reference number: I54011; MUID:88056305; PMID:3678832
A;Accession: I54011
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-72 <RE3>
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A;Cross-references: UNIPARC:UPI000016B0AE; GB:M17614; NID:g339483; PIDN:AAA61148.1; PID: C;Comment: Each of the two repetitive domains binds a ferric ion and a bicarbonate anion
 Proc. Natl. Acad. Sci. U.S.A. 86, 7260-7264, 1989 A;Title: Changes in brain gene expression shared by A;Reference number: I48174; MUID:89386721; PMID:2780 A;Accession: I63133
 A;Cross-references: UNIPARC:UPI000002F089; GB:S95936; N:R;Duguid, J.R.; Bohmont, C.W.; Liu, N.G.; Tourtellotte, Proc. Natl. Acad. Sci. U.S.A. 86, 7260-7264, 1989
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 A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-698 <RES>
 R;Hershberger, C.L.; Larson, J.L.; Arnold, B.; Rosteck, Ann. N. Y. Acad. Sci. 646, 140-154, 1991
A;Title: A cloned gene for human transferrin.
 Proc. Natl. Acad. Sci. U.S.A. 79, 2504-2508, 1982
A;Title: The complete amino acid sequence of human serum transferrin.
A;Reference number: A93911, MUID:82222166, PMID:6953407
A;Contents: annotation; disulfide bonds
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 A; Status: translated from GB/EMBL/DDBJ
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Best Local
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 Score 68; DB 1; ...
Pred. No. 0.0012;
 shared by scrapie and Alzheimer disease
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 DB 1; Length 698;
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 Woodworth,
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F;37-663/3757,136-212,155-349,176-192,179-195,109-197,245-259,357-611,363-395,373-386,420
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 17-Feb-1994 #sequence_revision 17-Feb-1994 #text_change 09-Jul-2004
C;Accession: A45543; S24173
C;Accession: A45543; S24173
Anim. Genet. 23, 251-256, 1992
A,Title: Cloning and sequencing of the porcine lactoferrin cDNA.
A,Reference number: A45543; MUID:92367939; PMID:1503259
 A;Note: 516-Ile was also found R;Heaphy, S.; Williams, J.
Blochem. J. 205, 611-617, 1982
A;Title: The preparation and partial characterization of N-terminal and C-terminal iron-k
A;Reference number: A26504; MUID:83074540; PMID:6816218
 A;Molecule type: protein
A;Residues: 482-515,'V',517-544 <EVA>
A;Cross-references: UNIPARC:UPI00001744A9
 R;Evans, R.W.; Aitken, A.; Patel, K.J.
FEBS Lett. 238, 39-42, 1988
A;Tille: Evidence for a single glycan moiety in rabbit serum transferrin
A;Reference number: S02694; MUID:89005676; PMID:3169252
 A;Molecule type: protein
A;Residues: 19-45,'S',47-48,'Y',50 <GOD>
A;Cross-references: UNIPARC:UPI00001744A8
 R;Godovac-Zimmermann, J.
Biol. Chem. Hoppe-Seyler 369, 93-96, 1988
A;Tille: Isolation, characterization and N-terminal amino-acid sequence of rabbit transfe A;Reference number: S00335; MUID:88209278; PMID:3365331
A;Accession: S00335
 A;Cross-references: UNIPARC:UPI00001744A6
R;Chung, M.C.M.; Chan, S.L.; Shimizu, S.
Int. J. Biochem. 23, 609-616, 1991
A;Title: Purification of transferrins and lactoferrin using A;Reference number: A61573; MUID:91293379; PMID:2065820
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A;Residues: 19-26,'X',28-36,'X',38-53 <CHU>
A;Cross-references: UNIPARC:UPI00001744A7
 A; Molecule type: protein
A; Residues: 19-36 < PIE>
 R;Pierpaoli, W.; Dall'Ara, A.; Yi, C.; Neri, P.; Santucci, A.; Choay, J. Cell. Immunol. 134, 225-234, 1991
A;Title: Iron carrier proteins facilitate engraftment of allogeneic bone A;Reference number: A61239; MUID:91191584; PMID:2013104
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 A; Accession: A26504
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2; Mismatches
 DB 1;
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A, Experimental source: manmary gland

A, Note: sequence excracted from NCBI backbone (NCBIN:111151, NCBIP:111153)

R; Lydon, J.P.; O'Malley, B.R.; Saucedo, O.; Lee, T.; Headon, D.R.; Conneely, O.M.

Biochim. Biophys. Acta 1132, 97-99, 1992

A; Title: Nucleotide and primary amino acid sequence of porcine lactoferrin.

A; Reference number: $24173; MUID:92379101; PMID:1511016

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F;359-696/Domain: transferrin repeat homology <TRH2>
F;252,300,387,495,564/Binding site: carbohydrate (Asn) (covalent)
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C;Accession: JC2323
 A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-703 <AL
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 Biochem. Biophys. Res. Commun. 203, 1324-1332, 1994
A;Title: Characterization of the goat lactoferrin cDNA:
A;Reference number: JC2323; MUID:94380047; PMID:8093048
A;Accession: JC2323
 C;Accession: JC2323

R;Le Provost, F.; Nocart, M.; Guerin, G.; Martin, P. Biochem. Biophys. Res. Commun. 203, 1324-1332, 1994
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 8
 transferrin precursor - rat
 S49163
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 A; Cross-references: UNIPROT: P14632; UNIPARC: UPI0000177C55
 RESULT 5
 lactoferrin - goat
 490/Binding site: carbohydrate
 Госат
 676
 1 YEKYLGEEYVKAV 13
 10;
 1-703 <ALE>
 Similarity
 Similarity
 YEKYLGTEYVTAI 688
 YEKYLGSEYVTAI 683
 YEKYLGEEYVKAV 13
 Conservative
 Conservative
 lung-derived growth factor; siderophilin
 80.9%;
 82.4%;
76.9%;
 Score 55; DB 2
Pred. No. 0.21;
1; Mismatches
 (Asn) (covalent) #status predicted
 Score 56; DB 2; Length 703; Pred. No. 0.14;
 Mismatches
12-May-1995 #text_change 09-Jul-2004
14679; A53289; A30512; I52203
 DB 2; Length 708;
 2
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 Indels
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 assignment of the relevant locu
 #status predicted
 0
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A; Note: sequence modified after extraction from NCBI backbone
A; Note: sequence extracted from NCBI backbone (NCBIP:86115, NCBIP:86116, NCBIP:86114)
R; Purves, L.R.; Purves, M.; Linton, N.; Brandt, W.; Johnson, G.; Jacobs, P.
Biochim. Biophys. Acta 966, 318-327, 1988
A; Title: Properties of the transferrin associated with rat intestinal mucosa.
A; Reference number: A30512; MUID:88327006; PMID:3046665
A; Molecula Francession: A30512
 A;Title: Transferrin messenger ribonucleic acid: molecular cloning and hormonal regulati A;Reference number: A30014; MUID:87053639; PMID:3023031
A;Recession: A30014
A;Molecule type: mRNA
A;Residues: 518-687,'D',689-692,'TA',695 <HUG-
A;Cross-references: UNIPARC:UPI0000170B4C; GB:M27966; NID:9207439; PIDN:AAA42267.1; PID: R;Schreiber, G.; Dryburgh, H.; Millership, A.; Matsuda, Y.; Inglis, A.; Phillips, J.; Ed A;Title: The synthesis and secretion of rat transferrin.
A;Reference number: A14679; MUID:80049855; PMID:500689
A;Accession: A14679
 A;Molecule type: protein
A;Residues: 20-30;639-643,'KD',646,'LKACD' <PUR>
A;Residues: 20-30;639-643,'KD',646,'LKACD' <PUR>
A;Cross-references: UNIPARC:UPI0000177C52; UNIPARC:UPI0000177C53
R;Aldred, A.R.; Howlett, G.J.; Schreiber, G.
Biochem. Biophys. Res. Commun. 122, 960-965, 1984
A;Title: Synthesis of rat transferrin in Escherichia coli containing a recombinant bacte
A;Reference number: 152203; MUID:84307580; PMID:6236811
 A;Molecule type: mRNA
A;Residues: 1-695 <ES2>
A;Residues: 1-695 <ES2>
A;Residues: 1-695 <ES2>
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 R;Escriva, H.; Pierce, A.; Coddeville, B.; Gonzalez, F.; Benaissa, M.; Leger, D.; Wieru Biochem. J. 307, 47-55, 1995
A;Title: Rat mammary-gland transferrin: nucleotide sequence, phylogenetic analysis and A;Reference number: S54980; MUID:95234054; PMID:7717992
A;Accession: S54980
 A;Molecule type: mRNA
A;Residues: 7-25,'X',27-56,'A',58-64,267-295 <RES>
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A;Residues: 20-47 <SCH>
A;Cross-references: UNIPARC:UPI0000177C51
A;Cross-references: UNIPARC:UPI0000177C51
Ccll. Biochem. 47, 261-271, 1991
J. Ccll. Biochem. 47, 261-271, 1991
A;Title: Lung-derived growth factor that stimulates the growth of lung-metastasizing
A;Reference number: A53289; MUID:92165927; PMID:1791188
 A_f Description: Rat mammary gland transferrin: glycan structure, A_f Reference \ number: S49163
 R;Escriva, H.; Pierce, A.; Coddeville, B.; Gonzalez, F.; Benaissa, M.; Leger, D.; Wierus submitted to the EMBL Data Library, January 1994
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 A;Status: preliminary
 A; Residues: 1-695 <ESC>
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transferrin repeat homology <TRH1>
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 PID:
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Similarity 9; Conserv

75.0%;

Score 51; D Pred. No. 1; Mismatches

DB 2;

Length 695; Indels

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A;Residues: 1-144,'V',146-163,'PP',166-339,'A',341-438,'Y',440-513,'R',515-708 <PII>A;Residues: 1-144,'V',146-163,'PP',166-339,'A',341-438,'Y',440-513,'R',515-708 <PII>A;Cross-references: UNIPARC:UPI00016C335; EMBL:X57084; NID:g505; PIDN:CAA40366.1; PIR:Cross-references: UNIPARC:UPI00016C335; EMBL:X57084; NID:g505; PIDN:CAA40366.1; PIR:Cross-references: Montreuil, J.; Spik, Sur. J. Biochem. 196, 177-184, 1991
A;Title: Molecular cloning and sequence analysis of bovine lactotransferrin.
A;Reference number: S14110; MUID:91160550; PMID:2001696
A;Accession: S14110
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A;Accession: S14110
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 carbonic anhydrase II inhibitor (transferrin homolog) precursor - pig (,Species: Sus scrofa domestica (domestic pig) (,Jate: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 09-Jul-2004 (,Accession: I47228 R.Roush, E.D., Fierke, C.A. Biochemistry 31, 12536-12542, 1992 A;Title: Purification and characterization of a carbonic anhydrase II inhib A;Reference number: I47228; MUID:93099129; PMID:1463741
 A;MOLECUle type: protein
A;Residues: 20-35;82-114;148-163,'PP',166-178,'V','P';183-190;205-212;230-239;304-339;59
A;Cross-references: UNIPARC:UPI000017448C; UNIPARC:UPI000017448D; UNIPARC:UPI000017448B;
493; UNIPARC:UPI0000174494
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C;Species: Bos primigenius taurus (cattle)
C;Date: 31-Mar-1992 #sequence revision 21-Nov-1997 #text change 09-Jul-2004
C;Accession: I45919; S14674; Si14110; S18517; JT0595; S13097; S18518; S13881;
R;Tsang, T.C.; Burns, D.K.; Wang, F.; Pan, Y.
RASEB J. 6, 233, 1991
A;Title: Cloning of a 80-kD advanced glycosylation end product (AGE) binding A;Reference number: I45919
 A;Gene: pICA
C;Superfamily: transferrin; transferrin repeat homology
C;Keywords: duplication
F;20-350/Domain: transferrin repeat homology <TRH1>
 A;Cross-references: UNIPROT:Q29545; C;Genetics:
 A; Molecule type: mRNA
A; Residues: 1-704 < ROU>
 A;Accession: I47228
A;Status: prelimina
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 submitted to the EMBL Data Library, November 1990 A;Reference number: $14674 A;Accession: $14674
 A;Cross-references: UNIPROT:P24627; UNIPARC:UPI0000137377; GB:L08604; NID:g163269; R;Pierce, A.
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 A;Residues:
 A; Accession: I45919
 N;Alternate names:
 actotransferrin
 Status: preliminary; translated from GB/EMBL/DDBJ
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 Query Match
Best Local
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 1-708 <TSA>
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 Similarity
8; Conserv
 YEEYLGAEYLQAV
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 YOKYLGPEYLOAI 684
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 Schanbacher, F.L
 Conservative
 lactoferrin
 75.0%;
 675
 13
180,
 4
 Score 51; I
Pred. No. 1;
75-84,
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 Mismatches
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 1;
 Length 704;
 Indels
 (AGE) binding protein
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 II inhibitor from
 Gaps
 PL0148;
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A;Cross references: UNIPARC:UPI00000475A8
R;Shimazaki, K.; Tanaka, T.; Kon, H.; Oota, K.; Kawaguchi, A.; Maki, Y.; Sato, T.
J. Dairy Sci. 76, 946-955, 1993
A;Title: Separation and characterization of the C-terminal half molecule of bovine A;Reference number: A56659; MUID:93253156; PMID:8486845
A;Accession: A56659
 F;79,111,211,272/Binding site: iron (Asp, Tyr, Tyr, His) #status F;140/Binding site: carbonate (Arg) #status experimental F;252,300,387,495,564/Binding site: carbohydrate (Asn) (covalent) F;414,452,545,614/Binding site: iron (Asp, Tyr, Tyr, His) #status
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A;Crose-references: UNIPARC:UPI00001744A0
R;Bellamy, W.; Takase, M.; Yamauchi, K.; Wakabayashi, H.; Kawase, Blochim. Biophys. Acta 1121, 130-136, 1992
A;Title: Identification of the bactericidal domain of lactoferrin.
A;Reference number: S21756; MUID:92287941; PMID:1599934
 submitted to the EMBL Data Library, October 1990 A;Reference number: S13881 A;Accession: S13881
 A;Residues: 20-47;59-66;132-139;256-277;278,305-332;343-351;361-363;586;587-589;598-619
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A;Crose-references: UNIPARC:UPI0000174495; GB:M63502
A;Note: the authors translated the codon CCG for residue 66
R;Mead, P.E.; Tweedle, J.W.
Nucleic Acids Res. 18, 7167, 1990
A;Title: cDNA and protein sequence of bovine lactoferrin.
A;Reference number: S13097; MUID:91088328; PMID:2263492
 A; Molecule type: protein
A; Residues: 36-60 < BEL>
 R;Rejman, J.J.; Hegarty, H.M.; Hurley, W.L.
Comp. Biochem. Physiol. B 93, 929-934, 1989
A;Tille: Purification and characterization of bovine lactoferrin
A;Reference number: PL0148; MUID:90031466; PMID:2805645
 A;Molecule type: mRNA
A;Residues: 28-38,'P',40-86,'C',88-708 <ME3>
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 A; Accession: S21756
 A;Accession: S18518
 A;Title: Bovine lactoferrin mRNA: sequence, analysis, and expression in the mammary gland
A;Reference number: JT0595; MUID:92028986; PMID:1718281
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 JT0595
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 YEKYLGEEYVKAV 13
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 Conservative
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 Length
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NyAlternate names: Iactorerrr...
C;Species: Homo sapiens (man)
C;Date: 31-Mar-1992 #sequence revision 21-Nov-1997 #text change 08-Dec-2000
C;Date: 31-Mar-1992 #sequence revision 21-Nov-1997 #text change 08-Dec-2000
 R;Gilmont, R.R.; Coulter, G.H.; Sylvester, S.R.; Griswold, M.D. Biol. Reprod. 43, 139-150, 1990
A;Title: Synthesis of transferrin and transferrin mRNA in bovine A;Reference number: A60166; MUID:90366647; PMID:2393886
A;Accession: A60166
 R;Stallard, B.J.; Collard, M.W.; Griswold, M.D. Mol. Cell. Biol. 11, 1448-1453, 1991
A;Title: A transferrinlike (hemiferrin) mRNA is expressed A;Reference number: A39684; MUID:91141493; PMID:1996102
A;Accession: A39684
 C;Species: Bos primigenius taurus (cattle)
C;Date: 03-Mar-1993 #sequence_revision 03-Mar-1993 #text_change 21-Nov-1997
C;Accession: A60166
 submitted to the EMBL Data A; Reference number: G06820 A; Accession: G01394
 A;Cross-references: UNIPROT:Q64599; UNIPARC:UPI00001744C3; C;Superfamily: hemiferrin; transferrin repeat homology C;Keywords: glycoprotein; testis
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C;Keywords: glycoprotein; testis
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 A;Cross-references: UNIPARC:UPI0000177C59
C;Comment: The authors designate this protein as transferrin. However, the closely resembles the C-terminal duplication in transferrin. This protein i
A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: mRNA
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 A; Molecule type: mRNA
A; Residues: 1-215 <GIL>
 A;Residues:
 A; Molecule type: mRNA
 hemiferrin -
 A; Status: not compared with conceptual translation
 hemiferrin -
 lactotransferrin precursor [validated] - human
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184 YDSYLGDDYVRAM 196
 183
 1-216 <STA>
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 Similarity
 bovine (fragment)
 YEKYLGEEYVKAV 13
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 transferrin repeat homology #status atypical <TRH2>
 Conservative
 Conservative
 70.6%;
53.8%;
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 Score 48; DB 1
Pred. No. 0.99;
 Score 48; DB 2;
Pred. No. 0.98;
5; Mismatches
 DB 1; Length 216;
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 A; Reference number: S74119; MUID: 97054624;
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A;Note: sequence extracted from NCBI backbone (NCBIP:122202) R;Powell, M.J.; Ogden, J.E. Nucleic Acids Res. 18, 4013, 1990 A;Title: Nucleotide sequence of human lactoferrin cDNA. A;Reference number: S10324; MUID:90326549; PMID:2374734 A;Accession: S10324
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 A,Experimental source: normal breast tissue R;Metz-Boutigue, M.H.; Jolles, J.; By Mazurier, J.; Schoentg Eur. J. Biochem. 145, 659-666, 1984 Eur. J. Biochem. 145, 659-666, 1984 Eur. A;Tille: Human lactotransferrin: amino acid sequence and A;Reference number: A31000; MUID:85076667; PMID:6510420 A;Accession: A31000
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R;Panella, T.J.; Liu, Y.; Huang, A.T.; Teng, C.T.
Cancer Res. 51, 3037-3043, 193
A;Title: Polymorphism and altered methylation of the lactoferrin gene in normal leukocyt
A;Title: Polymorphism and altered methylation of the lactoferrin gene in normal leukocyt
A;Reference number: A61169; MUID:91235214; PMID:1674448
 A;Cross-references: UNIPARC:UPI0000174487
R;Rado, T.A.; Wei, X.; Benz Jr., E.J.
Blood 70, 989-993, 197
A;Title: Isolation of lactoferrin cDNA from a human myeloid library and expression of A;Reference number: S07160; MUID:88001031; PMID:3477300
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A;Cross-references: UNIPARC:UPINOISC582; EMBL:X53961; NID:g34415; PIDN:CAA37914.1; PID R;Teng, C. T.; Liu, Y.; Yang, N.; Walmer, D.; Panella, T.

Mol. Endocrinol. 6, 1969-1981, 1992
A;Title: Differential molecular mechanism of the estrogen action that regulates lactofer A;Reference number: A45401; MUID:93125571; PMID:1480183
A;Accession: A45401
 A;Cross-references: UNIPARC:UPI000013737C; EMBL:U07643; NID:g467236; PIDN:AAB60324.1; PI R;Rey, M.W.; Woloshuk, S.L.; deBoer, H.A.; Pieper, F.R. Nucleic Acids Res. 18, 5288, 1990
Nucleic Acids Res. 18, 5288, 1990
A;Title: Complete nucleotide sequence of human mammary gland lactoferrin.
A;Reference number: S11228; MUID:90384839; PMID:2402455
 A;Title: Expression of cloned human lactoferrin in baby-hamster kidney cells A;Reference number: S15853; MUID:91264786; PMID:2049066 A;Accession: S15853
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A; Experimental source: gastric mucosa
A; Note: the authors suggest transferrin from gastric mucosa may act in dietary iron upta a; Diochem. 23, 609-616, 1991
A; Title: Purification of transferrins and lactoferrin using DEAE Affi-Gel Blue.
A; A; Cross-references: UNIPARC: UPI000174486
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C;Accession: S01384; A60520; A61573
R;Baldwin, G.S.; Weinstock, J.
Nucleic Acids Res. 16, 8720, 1988
A;Title: Nucleotide sequence of porcine liver transferrin.
A;Reference number: S01384; MUID:88335629; PMID:3419934
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A,Yoshinari, K.; Yuasa, K.; Iga, F.; Mimura, A.
Biochim. Biophys. Acta 1010, 28-34, 1989
A,Fitle: A growth-promoting factor for human myeloid leukemia cells from horse serum ider
A,Fitle: A growth-promoting factor for human myeloid leukemia cells from horse serum ider
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R;Liu, Y.; Teng, C.T.
J. Biol. Chem. 266, 21880-21885, 1991
A;Title: Characterization of estrogen-responsive mouse lactoferrin promoter.
A;Reference number: A41205; MUID:92042099; PMID:1939212
A;Accession: A41205
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R;Carpenter, M.A., Broad, T.E.
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A;Accession: S33761
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A;Title: Lactotransferrin is the major estrogen inducible protein of mouse uterine secret
A;Reference number: A92596; MUID:87280033; PMID:3611056
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C;Accession: A28438; A41205
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J. Bail. Chem. 262, 10134-10139, 1987
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A;Authors: Kaine, B.P.; Bordovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese, (A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii A;Reference number: A64300; MUID:96337999; PMID:8688087
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No. 0.023;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | ium; , Brandt P., Mewes H.W., Weil B., Amion M., Wiemann S.; the EMBL/GenBank/DDBJ databases. rot Consortium, see http://www.uniproireative Commons Attribution-NoDerivs ireative Common | ALIGNMENTS  LT 1  LT 1  LT PONPY  OFREL7,  OFREL7;  OFREL7;  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL7:  OFREL | 48 70.6 711 2 QSEK51 HUMAN  3 47 69.1 696 1 TREE PTG  4 46 67.6 309 2 Q5V6W4 HALMA  46 67.6 309 2 Q5V6W4 HALMA  47 66.7 6 647 1 DNAK PELUB  48 66.2 637 2 Q9TPQR4 HORSE  49 45 66.2 334 1 G3P PYRKO  49 45 66.2 334 1 G3P PYRKO  40 45 66.2 470 2 Q49ZP3 BLOPB  41 45 66.2 697 2 Q49ZP3 BLOPB  42 45 66.2 697 1 TREE MOUSE  43 46 6.2 697 2 Q3UBW7 MOUSE  44 66.2 697 2 Q3UBW7 MOUSE  45 66.2 706 1 TREE HORSE  46 66.2 707 1 TREE MOUSE  47 48 66.2 707 1 TREE MOUSE  48 66.2 707 1 TREE MOUSE  49 49 66.2 707 1 TREE MOUSE  40 49 66.2 707 1 TREE MOUSE  41 45 66.2 707 1 TREE MOUSE  42 45 66.2 707 1 TREE MOUSE  43 45 66.2 707 2 Q3TPZ4_MOUSE  44 66.2 707 2 Q3TPZ4_MOUSE  45 66.2 707 2 Q3TPZ4_MOUSE  46 66.2 707 2 Q3TPZ4_MOUSE |
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DT 24-MAY-2005, sequence version 1.
DT 24-MAY-2006, entry version 4.
DT 07-FEB-2006, entry version 4.
DE Transferrin variant (Fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebra
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 19-JUL-2005, integrated into UniProtKB/TrEMBL.
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Brain cDNA, clone: QccE-13766, similar to human transferrin (TF),.
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Cercopithecidae; Cercopithecinae; Macaca.
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Pfam; PF00405; Transferrin; 2.
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01-NOV-1990, integrated into UniProtKB/Swiss-Prot.
15-JUL-1999, sequence version 4.
07-MAR-2006, entry version 61.
Serotransferrin precursor (Transferrin) (Siderophilin) (Beta-1 metal-
 0000 SMR
 MEDLINE=91274362; PubMed=2054387; DOI=10.1016/0167-4781(91)90021-Banfield D.K., Chow B.K.-C., Funk W.D., Robertson K.A., Umelas T. Woodworth R.C., Macgillivray R.T.A.; "The nucleotide sequence of rabbit liver transferrin cDNA."; Biochim. Biophys. Acta 1089:262-265(1991).
 Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporid
 Name=TF
 PRINTS; PRO0422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00206; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 2; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
SEQUENCE 698 AA; 77010 MW; F19A15A7C898399B
 GO; GO:0005576; C:extracellular region; IEA
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0008679; P:iron ion homeostasis; IEA
GO; GO:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11455; Peptidase_S60; 1.
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 evolutionary implications.";
Submitted (JUN-2005) to the EMBL/GenBank/DDBJ
 "Sugano S., Gojobori T., Shen C.-K.J., Wu C.I., Has "Substitution Rate and Structural Divergence of S' Comparative Analysis Between Human and Cynomolgus Mol. Biol. Evol. 22:1976-1982(2005).
 "Cloning and structural organisation encoding gene.";
 Ghareeb B.A.A., Houdebine L.M.;
 binding globulin).
 EMBL;
 PROTEIN SEQUENCE OF 20-51
MEDLINE=88209278; PubMed=
 STRAIN=New
 NUCLEOTIDE SEQUENCE [MRNA].
 ; AB169522; BAE01604.1; -; mRNA.
Q4R5M1; 23-346.
 sequences of macaque
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 InterPro; IPRO01156; Peptidase_S60.

PANTHER; PTHR11485; Peptidase_S60; 1.

Pfam; PF00405; Transferrin; 2.

PRINTS; PR00422; TRANSFERRIN.

SMART; SM00094; TR FER; 2.

PROSITE; PS00205; TRANSFERRIN 1; 1.

PROSITE; PS00205; TRANSFERRIN 2; 2.

PROSITE; PS00206; TRANSFERRIN 3; 2.

PROSITE; PS00207; TRANSFERRIN 3; 2.
 EMBL;
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 -!- SUBUNIT: Monomer.
-!- SUBCELLULAR LOCATION: Secreted protein.
-!- TISSUE SPECIFICITY: Expressed by the liver and
-!- TISSUE SPECIFICITY: Expressed by the liver and
-!- MISCELLANEOUS: The peptidase $60 domains do not
function as the catalytic residues are missing
-!- SIMILARITY: Belongs to the transferrin family.
-!- SIMILARITY: Contains 2 peptidase $60 domains.
 MEDLINE=89026775; PubMed=3179277;
Bailey S., Evans R.W., Garratt R.C., (
Horsburgh C., Jhoti H., Lindley P.F.,
"Molecular structure of serum transfe:
Biochemistry 27:5804-5812(1988).
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 X-RAY CRYSTALLOGRAPHY (2.3 ANGSTROMS).
Sarra R., Garratt R.C., Gorinsky B., Jhoti H., Lindley P.F.;
"High-resolution X-ray studies on rabbit serum transferrin:
preliminary structure analysis of the N-terminal half-molecule
 MEDLINE=89005676; PubMed=3169252; DOI=10.1016/0014-5793(88)80221-7; Evans R.W., Aitken A., Patel K.J.; "Evidence for a single glycan moiety in rabbit serum transferrin and location of the glycan within the polypeptide chain."; TEBS Lett. 238:39-42(1988).
 MEROPS; $60.972; -. MEROPS; $60.975; -.
 EMBL; AF031624; APPDB; LJNF; X-ray; PDB; LTFD; X-ray;
 EMBL;
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
 "Isolation, characterization and N-terminal rabbit transferrin.";
 LinkHub; P19134;
 TISSUE=Plasma;
 X-RAY CRYSTALLOGRAPHY (3.3
 PROTEIN SEQUENCE OF
 Godovac-Zimmermann J.;
 resolution."
 FUNCTION: Transferrins are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate. It is responsible for the transport of iron from sites of absorption and heme degradation to those of storage and utilization. Serum transferrin may also have a further role in stimulating cell proliferation.
 Crystallogr. B 46:763-771(1990) FUNCTION: Transferrins are iron
 AF031616;
AF031617;
AF031618;
AF031619;
AF031620;
 Chem. Hoppe-Seyler 369:93-96(1988)
 AF031625; AAB94136.1;
 X58533; CAA41424.1;
 AF031615;
 AF031611;
 AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; JOINED; AAB94136.1; AAB94136.1; JOINED; AAB94136.1; AA
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GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion themeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
Interpro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
PFAm; PF00405; Transferrin; 1.
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NCBI_TaxID=9823;
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Shinkai H., Morozumi T., Toki D., Muneta Y., Awata T., U Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
 Sus scrofa (Pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suina; Suidae;
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82.4%; Score 56;
 Score 58; I
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P14632; Q29557;
01-APR-1990, integrated into UniProtKB/Swiss-Prot.
15-DEC-1998, sequence version 3.
07-FEB-2006, entry version 46.
Lactotransferrin precursor (EC 3.4.21.-) (Lactofer
NUCLEOTIDE SEQUENCE [MRNA]:
NUCLEOTIDE SEQUENCE [MRNA]:
MEDLINE=92379101; PubMed=1511016; DOI=10.1016/0167-4781(92)90061-4;
MEDLINE=92379101; PubMed=1511016; DOI=10.1016/0167-4781(92)90061-4;
 Sus scrofa (Pig).
Eukaryota; Metazoa;
 PRINTS; PR00422; TRANSFERRIN.
SMART; SM00094; TR FER; 2.
PROSITE; PS00205; TRANSFERRIN 1; 2.
PROSITE; PS00206; TRANSFERRIN 2; 2.
PROSITE; PS00207; TRANSFERRIN 3; 2.
SEQUENCE 694 AA; 76466 MW; 4005
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GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; F:iron ion homeostasis; IEA
GO; GO:0006826; F:iron ion transport; IEA.
InterPro; IFR001156; Peptidase S60.
PANTHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 2.
 Name=LTF
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Rinaudo J.A.S., Gerin J.L.;
Submitted (APR-2003) to the EMBL/GenBank/DDBJ
 Mammalia; Eutheria;
Sciuridae; Xerinae;
 Eukaryota; Metazoa;
 01-OCT-2003, sequence version 1.
07-FEB-2006, entry version 11.
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 Mammalia; Eutheria;
 NCBI_TaxID=9995;
 Marmota monax (Woodchuck)
 Transferrin.
 01-OCT-2003,
 NUCLEOTIDE SEQUENCE
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 SSUE=Liver;
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 P19134; 1JNF.
 AY288100; AAP37129.1; -; mRNA.
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(EC 3.4.21.-) (Lactoferrin)
 Score 56; DB
Pred. No. 2.3;
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PROSITE; PS00205; TRANSFERRIN_2; 2.
PROSITE; PS00206; TRANSFERRIN_3; 1.
PROSITE; PS00207; TRANSFERRIN_3; 1.
 Hutchens T.W., Magnuson J.S., Yip T.-T.;
"Rapid purification of porcine colostral whey lactoferrin by affichromatography on single-stranded DNA-agarose. Characterization, acid composition and N-terminal amino acid sequence.";
Biochim. Biophys. Acta 999:323-329(1989).
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 EMBL; M92089; AAA31102.1; -;
EMBL; M81327; AAA31059.1; -;
PIR; A45543; A45543.
HSSP; Q9TUMO; 1DTZ.
 Anim.
 NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=92367939; PubMed=1503259;

Alexander L.J., Levine W.B., Teng C.T., Beattie C.W.;

"Cloning and sequencing of the porcine lactoferrin cDNA.";

Anim. Genet. 23:251-256(1992).
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 Serine protease; Signal; SIGNAL 1 19
 Direct
 InterPro; IPR001156; Peptidase_S60. PANTHER; PTHR11485; Peptidase_S60; Pfam; PF00405; Transferrin; 2.
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 -
 PROTEIN SEQUENCE OF 20-49.
MEDLINE=90105538; PubMed=2605266;
 Biochim. Biophys. [2]
 CHAIN
 Iron;
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 Conneely O.M.
 "Nucleotide
 SUBCELLULAR LOCATION: Secreted protein.
MISCELLANEOUS: The second peptidase S60 domain does not protease function as the catalytic residues are missing.
SIMILARITY: Belongs to the transferrin family.
SIMILARITY: Contains 2 peptidase S60 domains.
 FUNCTION: Transferrins are iron binding transport proteins which can bind two Fe(3+) ions in association with the binding of an anion, usually bicarbonate.

FUNCTION: The lactotransferrin peptidase S60 domain 1 functions as a serine protease that cuts arginine rich regions. This function contributes to the antimicrobial activity (By similarity).

CATALYTIC ACTIVITY: Preferential at -Arg-Ser-Arg-|- and -Arg-Arg-Ser-Arg-|- and of z-Phe-Arg-|-aminomethylcoumarin.

SUBUNIT: Monomer
 t protein sequencing; Glycoprotein; Hydrolase; Ion transport;
Iron transport; Metal-binding; Protease; Repeat;
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05-JUL-2004, sequence vers
07-FEB-2006, entry version
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 BINDING
 Distributed under the Creative Commons Attribution-NoDerivs
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 Eukaryota; Metazoa;
 Sus scrofa (Pig)
 SEQUENCE
 CARBOHYD
 BINDING
 NUCLEOTIDE SEQUENCE.
 NCBI_TaxID=9823;
 Name=LTF
 Lactotransferrin.
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 P; Q29477; iJW1.
GO:0005576; C:extracellular region; IEA.
GO:0008199; F:ferric iron binding; IEA.
GO:0006879; P:iron ion homeostasis; IEA.
GO:0006876; P:iron ion transport; IEA.
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 YEKYLGSEYVTAI
 IPR001156; Peptidase_S60
 Eutheria;
 704 AA;
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 integrated into UniProtKB/TrEMBL.
 PRELIMINARY;
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GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
 InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
Pfam; PF00405; Transferrin; 2.
 TISSUE=Mammary gland;
Pecorini C., Fogher C., Baldi A.;
Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suina; Suidae;
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 NCBI_TaxID=9823;
 Sus scrofa (Pig).
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 PRINTS; PRO0405; Transferrin; 2

PRINTS; PRO0422; TRANSFERRIN.

PROSITE; PRO0205; TRANSFERRIN.

PROSITE; PRO0205; TRANSFERRIN.

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 HSSP; P24627; INKX.

HSSP; P24627; INKX.

GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006879; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase S60.
PANTHER; PTHR11485; Peptidase S60; 1.
 TISSUE-Mammary gland;
TISSUE-Mammary gland;
MEDLINE-94380047; PubMed-8093048;
MEDLINE-94380047; PubMed-8093048;
le Provost F., Nocart M., Guerin G., Martin P.;
le Provost F., Nocart M., Guerin G., Martin P.;
Characterization of the goat lactoferrin cDNA. Assignm relevant locus to bovine Ul2 synteny group.";
Plochem. Biophys. Res. Commun. 203:1324-1332(1994).
-I- FUNCTION: Transferrins are iron binding transport.
-I- FUNCTION: Transferrins are iron binding transport.
 Name=LTF;
Sus scrofa (Pig).
Eukaryota, Metazoa; (
Mammalia; Eutheria;)
 TREL CAPHI STANDARD; PRT; 708 AA. 029477; Q29479; 15-DEC-1998, integrated into UniProtKB/Swiss-Prot. 01-NOV-1996, sequence version 1. 07-FEB-2006, entry version 41. Lactotransferrin precursor (EC 3.4.21.-) (Lactofer
 TISSUE=Mammary gland;
Lee T., Yu S., Kim S., Lee K.,
Submitted (APR-1996) to the EM
 Mammalia; Eutheria; Lauras
Pecora; Bovidae; Caprinae;
 Capra hircus (Goat).
Eukaryota, Metazoa,
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 TISSUE=Mammary gland; Wang S.-R., Lin T.-Y., Weng Submitted (DEC-2001) to the
 -
 Name=LTF;
 EMBL; L77887; AAL40161.1; -; mRNA.
 NCBI_TaxID=9823;
 NUCLEOTIDE
 can bind two Fe(3+) ions in association with the bir anion, usually bicarbonate.
FUNCTION: The lactotransferrin peptidase S60 domain a serine protease that cure avainable.
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 YEKYLGSEYVTAI 684
 SEQUENCE
 Metazoa;
 SEQUENCE
 Conservative
 Laurasiatheria;
prinae; Capra.
 Chordata; Craniata; Vertebrata; Euteleostomi; Laurasiatheria; Cetartiodactyla; Ruminantia;
 Chordata; Craniata; Vertebrata; Euteleostomi; Laurasiatheria; Cetartiodactyla; Suina; Suidae;
 ., Lee K., Yu D.;
to the EMBL/GenBank/DDBJ databases
 [MRNA].
 82.4%;
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EMBL/GenBank/DDBJ databases.
 Score 56; 1
Pred. No. 2
 3.4.21.-) (Lactoferrin)
 64EE769F7503CC32 CRC64;
 ed. No. 2.3;
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MEROPS; SOLUCIAN MEROPS; SOLUCIAN MEROPS; INDOING PARTHER; PTHRI1485; Peptidase S60; 1.

PANTHER; PTHRI1485; Peptidase S60; 1.

Pfam; PF00405; Transferrin. 2.

PRINTS; PR00422; TRANSFERRIN. 1; 2.

PROSITE; PS00205; TRANSFERRIN. 1; 2.

PROSITE; PS00206; TRANSFERRIN. 2; 2.

PROSITE; PS00207; TRANSFERRIN. 3; 2.

PROSITE; PS00207; TRANSFERRIN. 3; 2.

PROSITE; PS00207; TRANSFERRIN. 3; 2.
 Iron transport; Metal-binding; Signal; Transport. SIGNAL 19 By
 EMBL; U53857; AAA97958.1; -; EMBL; X78902; CAA55517.1; -; PDB; 1JW1; X-ray; A=20-708.
 CONTRIBUTES to the antimicrobial activity (By similarity). CATALYTIC ACTIVITY: Preferential at -Arg-Ser-Arg-Arg-|- at Arg-Ser-Arg-|-, and of Z-Phe-Arg-|- aminomethylcoumarin. SUBUNIT: Monomet (By similarity).

SUBCELLULAR LOCATION: Secreted protein.
MISCELLANBOUS: The second peptidase $60 domain does not he protease function as the catalytic residues are missing. SIMILARITY: Belongs to the transferrin family.

SIMILARITY: Contains 2 peptidase $60 domains.
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Matches 10
 EMBL; AY792499; AAV92908.1; -; mRNA.

SMR; OSMJEB; 22-708.

GO; GO:0005576; C:extracellular region; IEA.

GO; GO:0005576; C:extracellular region; IEA.

GO; GO:0008199; F:ferric iron binding; IEA.

GO; GO:0006879; P:iron ion homeostasis; IEA.

GO; GO:0006879; P:iron ion transport; IEA.

InterPro; IPR001156; Peptidase S60.

PANTHER; PTHR11485; Peptidase S60; 1.

Pfam; PF00405; Transferrin; 2.

PRINTS; PR00422; TRANSFERRIN.

SMART; SM00094; TR FER; 2.

PROSITE; PS000205; TRANSFERRIN—1; 2.

PROSITE; PS000205; TRANSFERRIN—2; 2.

PROSITE; PS000207; TRANSFERRIN—3; 2.

SEQUENCE 708 AA; 77236 MW; BOAEEEFC69BBA2CC
 JULT 13
JUES SHEEP PRELIMINARY; PRT;
Q5MJES;
Q5MLES;
Q1-FEB-2005, integrated into UniPro
Q1-FEB-2005, sequence version 1.
Q7-FEB-2006, entry version 5.
 Ovis aries (Sheep).

Gukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;

Pecora; Bovidae; Caprinae; Ovis.

NCBI TaxID=9940;
 TISSUE=Mammary gland;
Grover S., Geetha N., Mohanty A.K., Kitamura Y., Batish
Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 NUCLEOTIDE SEQUENCE
 Lactoferrin.
 676
 10;
 1 YEKYLGEEYVKAV 13
 Similarity 76.9
LO; Conservative
 YEKYLGTEYVTAI 688
 integrated into UniProtKB/TrEMBL
 3305788
33057917
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 80.9%;
 Score 55; DB
Pred. No. 3.4;
1; Mismatches
 PRT;
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 708
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DT 05-JUL-2004, integrated into UniProtKB/TrEMBL.
DT 05-JUL-2004, sequence version 1.
DT 07-FEB-2006, entry version 10.
DT Lactoferrin (Fragment).
OS Bos taurus (Bovine).
OC Eukaryotta; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
OC Pecora; Bovidae; Bovinae; Bos.
OX NCBI_TaxID=9913;
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 Query Match
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 Ensembl; ENSETAG0000001292; Bos taurus.
G0; G0:0005576; C:extracellular region; IEA.
G0; G0:0008199; F:ferric iron binding; IEA.
G0; G0:00068199; F:iron ion homeostasis; IEA.
G0; G0:0006879; P:iron ion transport; IEA.
G0; G0:0006826; P:iron ion transport; IEA.
InterPro; IPR001156; Peptidase_S60.
PANTHER; PTHR11485; Peptidase_S60; 1.
 EMBL; X17066; CAA34912.1; -; mRNA.
HSSP; P24627; 1BLF.
SMR; Q8MIIO; 8-347.
 Pierce A., Colavizza D., Benaissa M., Maes P., Tartar A., Montreuil J., Spik G.; Montreuil J., Spik G.; "Molecular cloning and sequence analysis of bovine lactotransferrin."; Eur. J. Biochem. 196:177-184(1991).
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
Pecora; Bovidae; Bovinae; Bos.
 07-FEB-2006, entry version 14. Lactotransferrin (Fragment).
 QBMIIO BOVIN PRELIMINARY; PRT; 355 AA. Q8MIIO; 01-QCT-2002, integrated into UniProtKB/TrEMBL
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 MEDLINE-91160550; PubMed=2001696;
Pierce A., Colavizza D., Benaissa M., Maes P., Tartar A.,
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 Bos taurus (Bovine).
 01-OCT-2002, sequence version 1.
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 YEKYLGEEYVKAV 13
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 76.9%;
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 Length 355;
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SEQUENCE
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GO; GO:000819; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006826; P:iron ion transport; IEA.
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PANTHER; PTHR11485; Peptidase S60; 1.
Pfam; PF00405; Transferrin; 2.
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HSSP; Q29477; 1JW1.
SMR; Q6LEN7; 1-681.
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MEDLINE=9108838; PubMed=2263492;
MEDLINE=9108838; PubMed=2263492;
Mead P.E., Tweedie J.W.;
"CDNA and protein sequence of bovine lactoflavin.";
Nucleic Acids Res. 18:7167-7167(1990).
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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Submitted (JAN-1991) to the EMBL/GenBank/DDBJ databases
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 (1)
NUCLEOTIDE SEQUENCE.
 649
 1 YEKYLGEEYVKAV 13
 9; Conserv
 YEEYLGTEYVTAI 661
 681 AA; 75182 MW; 4927DE88144D99BA CRC64;
 Conservative
 75.0%;
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Pred. No.
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 DB 2;
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 Length 681;
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 Minimum DB seq
Maximum DB seq
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 Total number of hits satisfying chosen parameters:
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 Sequence:
 Perfect score:
 Title:
 Score
 d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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 US-10-612-162A-4
68
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1074
1074
11074
1410
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1703
703
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 GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd
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 US-09-439-740-2
US-08-470-058-2
US-09-285-310-2
US-09-285-310-3
US-09-285-310-4
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 Patent No
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RESULT 1
5262177-4
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 US-08-175-158A-2
 5262177-4
 SEQ ID NO:4:
 Matches
 Query Match
Best Local
 STATE: M
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|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|
| 48              | 48              | 48              | 48              | <b>4</b> 8      | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              | 48              |
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| 711             | 711             | 711             | 711             | 711             | 711             | 711             | 711             | 711             | 711             | 711             | 709             | 709             | 709             | 709             | 709             | 708             | 705             | 694             |
| N               | N               | N               | N               | N               | ۳               | ۳               | -ر              | _               | ۳               | ۲               | N               | N               | N               | ۳               | <b>ب</b>        | _               | <b>-</b>        | N               |
| US-09-265-577-2 | US-08-476-798-4 | US-08-456-108-2 | US-09-158-313-4 | US-08-464-167-4 | US-08-456-106-2 | US-08-453-703-2 | US-08-461-333-4 | US-08-154-019-4 | US-08-250-308-2 | US-08-145-681-2 | US-08-476-798-2 | US-09-158-313-2 | US-08-464-167-2 | US-08-461-333-2 | US-08-154-019-2 | US-08-655-640-4 | US-08-655-640-2 | US-09-932-190-2 |
| Sequence 2,     | Sequence 4,     | Sequence 2,     | Sequence 4,     | Sequence 4,     | Sequence 2,     | Sequence 2,     | Sequence 4,     | Sequence 4,     | Sequence 2,     | Sequence 2,     | Sequence 2,     | Sequence 2,     | Sequence 2,     | Sequence 2,     | Sequence 2,     | Sequence 4,     | Sequence 2,     | Sequence 2,     |
| , Appli         | , Appli         | , Appli         | , Appli         | , Appli         | Appli           | Appli           | Appli           | , Appli         | Appli           | , Appli         | , Appli         | Appli           | , Appli         | , Appli         | , Appli         | , Appli         | Appli           | Appli           |

ALIGNMENTS

```
; PATENT NO. 5262177
; PATENT NO. 5262177
; APPLICANT: BROWN, J OSEPH P.; ESTIN, CHARLES D.; PLOWMAN, GREGORY
; D.; HELLSTROM, KARL E.; ROSE, TIMOTHY M.; HELLSTROM, INGEGERD;
; PURCHIO, ANTHONY F.; HJ, SHIU-LOX; PENNATHUR, SRIDHAR
; TITLE OF INVENTION: RECOMBINANT VIRUSES ENCODING THE HHUMAN
; MELANOWA-ASSOCIATED ANTIGEN
; MUMBER OF SEQUENCES: 6
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/7230
; PRIOR APPLICATION DATA:
; APPLICATION DATA:
; APPLICATION NUMBER: 827,313
; PRIOR APPLICATION NUMBER: 827,313
; PRIOR APPLICATION NUMBER: 827,313
 Sequence 2, Applic
Patent No. 5986067
 GENERAL INFORMATION:
APPLICANT: FUNK, Walter D.
APPLICANT: MAGGILLIVRAY, Ross T.A.
APPLICANT: MASON, Anne B.
APPLICANT: MOODWORTH, Robert C.
TITLE OF INVENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
TITLE OF INVENTION: MOLECULES AND MUTANTS THEREOF
NUMBER OF SEQUENCES: 7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 CORRESPONDENCE ADDRESS:
 ADDRESSEE:
STREET: 60
 LENGTH: 696
 664 YEKYLGEEYVKAV 676
 ch 100.0%; Score 68; DB 7; Length 696; l Similarity 100.0%; Pred. No. 0.0012; 13; Conservative 0; Mismatches 0; Indels
 1 YEKYLGEEYVKAV 13
 Boston
 02109
 Massachusetts
 Application US/08175158A
 60 State Street,
 USA
 LAHIVE & COCKFIELD
 suite
 510
 0,
 Gaps
```

OPERATING SYSTEM: PC-D SOFTWARE: ASCII text CURRENT APPLICATION DATA:

COMPUTER:

IBM PC compatible

PC-DOS/MS-DOS

US/08/175,158A

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 ; MOLECULE TYPE: protein US-08-175-158A-2
 REFERENCE/DOCKET NUMBER: 31,503
REFERENCE/DOCKET NUMBER: UVI-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 698 amino acids
TYPE: amino acid
TOPOLOGY: linear
 US-09-439-740-2
 Patent No. 682503
 Matches
 Query Match
Best Local Similarity
 ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/175,158A FILING DATE: 28-DEC-1993 CLASSIFICATION: 530 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07/832,029 FILING DATE: 06-FEB-1992 ATTORNEY/AGENT INFORMATION:
 APPLICANT: MASON, APPLICANT: MASON, APPLICANT: WOODWORTH, ROBERT C.
APPLICANT: WOODWORTH, RECOMBINANT TRANSFERRINS, INTELLE OF INVENTION: MOLECULES AND MUTANTS THEREOF NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE LAHIVE & COCKFIELD
CORRESSEE: LAHIVE & COCKFIELD
 GENERAL INFORMATION:
 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/175,158
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: DeConti, Gillio A.
REGISTRATION NUMBER: 31,503
REFERENCE/DOCKET NUMBER: UVI-005CP2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
 APPLICANT:
 CITY: Boston
STATE: Massachusetts
COUNTRY: USA
 APPLICATION NUMBER: US/09/439,740 FILING DATE:
 NAME: DeConti, Giulio REGISTRATION NUMBER: 3
 666 YEKYLGEEYVKÁV 678
 1 YEKYLGEEYVKAV 13
 Application US/09439740
 MASON, Anne B.
WOODWORTH, Robert C.
WENTION: RECOMBINANT TRANSFERRINS, TRANSFERRIN HALF-
 FUNK, Walter D.
MacGILLIVRAY, Ross T.A.
 100.0%; Score 68; DB 1; nilarity 100.0%; Pred. No. 0.0012; Conservative 0; Mismatches 0
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US-09-037-188-2
; Sequence 2, Application US/09037188
; Patent No. 607921
; GENERAL INFORMATION:

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ZIP: 02173
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/08/470,058
FILING DATE: 06-JUN-1995
CLASSIFICATION: 1995
CLASSIFICATION: NOWNER: TS0
NAME: Granahan, Patricia
REFINATION NUMBER: TXT93-01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
2:
 ; MOLECULE TYPE: protein US-08-470-058-2
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 US-08-470-058-2
 US-09-439-740-2
 Query Match
Best Local Similarity
Matches 13; Conserv
 Sequence 2, Application US/08470058 Patent No. 5817789 GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 INFORMATION FOR SEQ ID NO:
 APPLICANT: Heartlein, Michaei m.
APPLICANT: Lemontt, Jeffrey F.
TITLE OF INVENTION: Chimeric Proteins For Use in Transport
TITLE OF INVENTION: of a Selected Substance Into Cells
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
 SEQUENCE CHARACTERISTICS:
LENGTH: 698 amino acids
TYPE: amino acid
TYPE: amino acid
 SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino acids
 TOPOLOGY: linear MOLECULE TYPE: protein
 TYPE: amino acid
STRANDEDNESS: unkr
TOPOLOGY: unknown
 STREET: Two Mili
 ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C STREET: Two Militia Drive
1042 YEKYLGEEYVKAV 1054
 666 YEKYLGEEYVKÁV 678
 13; Conservative
 1 YEKYLGEEYVKAV 13
 1 YEKYLGEEYVKAV 13
 100.0%; Score 68; DB 1; ilarity 100.0%; Pred. No. 0.0019; Conservative 0; Mismatches
 unknown
 100.0%; Score 68; DB 2; 100.0%; Pred. No. 0.0012;
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 Gaps
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 COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 06-UNN-1995
FILING DATE: 06-UNN-1995
 GENERAL INFORMATION:
 TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino aci
 NAME: Fraser, Ph.D., J.D., Ja
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 0723
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
 APPLICANT: Heartlein, Michael W.
APPLICANT: Lemontt, Jeffrey F.
APPLICANT: Concino, Michael F.
TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE
TITLE OF INVENTION: IN TRANSPORT OF A SELECTED
NUMBER OF SEQUENCES: 12
 MOLECULE TYPE: protein
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
 TITLE OF INVENTION:
TITLE OF INVENTION:
 ATTORNEY/AGENT INFORMATION:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
 CORRESPONDENCE ADDRESS:
 STATE:
 TOPOLOGY: unknown
 OPERATING SYSTEM:
SOFTWARE: FastSE(
 COUNTRY: USA
ZIP: 02110-2804
 ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street
 ZIP: 02110-2804
 COUNTRY:
 1042 YEKYLGEEYVKAV 1054
 13;
 1 YEKYLGEEYVKAV 13
 Boston
 Boston
 ¥
 3
 1074 amino acids
 Conservative
 USA
 Lemontt, Jeffrey F.
Concino, Michael F.
CENTION: CHIMERIC PROTEINS FOR USE
VENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
 Heartlein, Michael W.
 IBM Compatible
 Diskette
 Windows95
 0
 07236/009002
 Janis K.
 Mismatches
 SELECTED SUBSTANCE INTO CELLS
 Length 1074;
 0;
 Gaps
 0
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```
APPLICANT: Hemontt, Jeffrey F.

APPLICANT: Lemontt, Jeffrey F.

APPLICANT: Concino, Michael F.

APPLICANT: Concino, Michael F.

APPLICANT: Concino, Michael F.

APPLICANT: Concino, Michael F.

TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS

FILE REFERENCE: 10278-025004

CURRENT APPLICATION NUMBER: US/09/753,385

CURRENT APPLICATION NUMBER: US 08/470,058

PRIOR APPLICATION NUMBER: US 08/470,058

PRIOR FILING DATE: 1998-03-09

PRIOR FILING DATE: 1998-03-09

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR FILING DATE: 1998-03-09

PRIOR APPLICATION NUMBER: US 09/285,310

PRIOR FILING DATE: 1999-04-02
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RESULT 8
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 ; ORGANISM: Homo sapiens
US-09-753-385-2
 US-09-753-385-2
 US-09-285-310-2
 NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 1074
 Sequence 2, Application US/09753385 Patent No. 6858578 GENERAL INFORMATION:
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 INFORMATION FOR SEQ ID NO:
 TYPE: PRT
 NAME: Fraser, Ph.D., J.D., V. REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07.
TELECOMMUNICATION INFORMATION:
TELECHONE: 617,542-5070
 SEQUENCE CHARACTERISTICS:
LENGTH: 1074 amino acids
 TOPOLOGY: unknown MOLECULE TYPE: protein
 FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Ph.D., J.D
 FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
 Local Similarity 100.0%;
nes 13; Conservative (
 APPLICATION NUMBER: US/09/285,310
 TELEFAX:
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 1042 YEKYLGEEYVKAV 1054
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)GY: unknown
 617/542-8906
 Conservative
 100.0%; Score 68; 100.0%; Pred. No.
 09/037,188
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 Janis K.
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 DB 2;
 Length 1074;
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 Gaps
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US-08-470-058-4

Sequence 4, Application US/08470058 Patent No. 5817789

GENERAL INFORMATION:
APPLICANT: Heartle
APPLICANT: Lemont

Heartlein, Michael W. Lemontt, Jeffrey F.

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 US-09-037-188-4
 RESULT 9
 ; TOPOLOGY: unknown ; MOLECULE TYPE: protein US-08-470-058-4
 Sequence 4, Application US/09037188 Patent No. 6027921
 Matches
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Best Local Similarity
 TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/470,058 FILING DATE: 06-JUN-1995 CLASSIFICATION: 530
 COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/037,188
FILING DATE: 02-MAR-1998
 COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
 NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: TK:
TELECOMMUNICATION INFORMATION:
TELECHOONE: 617-861-6240
 TITLE OF INVENTION: Chimeric Proteins For Use in TraTITLE OF INVENTION: of a Selected Substance Into Cel NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
 FILING DATE: 02-MAR-: PRIOR APPLICATION DATA: APPLICATION NUMBER:
 TITLE OF INVENTION:
 APPLICANT:
 ATTORNEY/AGENT INFORMATION:
 COMPUTER READABLE FORM:
 APPLICANT:
 STREET: 1*C
 NUMBER OF SEQUENCES:
 TYPE: amino acid
STRANDEDNESS: unkr
TOPOLOGY: unknown
 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
 COUNTRY: USA
ZIP: 02173
 LENGTH:
 1378 YEKYLGEEYVKAV 1390
 INFORMATION:
 13;
 1 YEKYLGEEYVKAV 13
 Boston
 ₹
 1410 amino acids
 Lemontt, Jeffrey F.
Concino, Michael F.
TVENTION: CHIMERIC PROTEINS FOR USE
TVENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
 100.0%; Score 68; DB 1; Length 1410; nilarity 100.0%; Pred. No. 0.0026; Conservative 0; Mismatches 0; Indels
 Heartlein, Michael W.
06-JUN-1995
 Floppy disk
 unknown
 Chimeric Proteins For Use in Transport of a Selected Substance Into Cells 12
 TKT93-01
 0
 Gaps
 0
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Query Match
Best Local Similarity
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 ; MOLECULE TYPE: protein US-09-285-310-4
 밁
 US-09-037-188-4
 US-09-285-310-4
 Patent No. 6262026
GENERAL INFORMATION:
APPLICANT: Heartle:
APPLICANT: Lemontt
APPLICANT: Concino
 IELEX: 20154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1410 amin
 Matches
 Best Local
 Query Match
 INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 1410 amino acids
 COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
 NAME: Fraser, Ph.D., J.D., Janis K. REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 07236/009002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/037,188
FILLING DATE:
ATTORNEY/AGENT INFORMATION:
 ATTORNEY/AGENT INFORMATION:
NAME: FRABER, Ph.D., J.D., CANTON OF THE PROJECT AUMBER: 34,819
REFERENCE/DOCKET NUMBER: 077
 NUMBER OF SEQUENCES: 1
 TITLE OF INVENTION:
 MOLECULE TYPE: protein
 TELEPHONE: 617/542-5070
 STREET: 225 F
CITY: Boston
STATE: MA
 TOPOLOGY:
 TELEFAX:
 APPLICATION NUMBER: FILING DATE:
 TOPOLOGY:
 ADDRESSEE:
 1378 YEKYLGEEYVKAV 1390
 4.
 1 YEKYLGEEYVKAV 13
 Similarity
 amino acid
GY: unknown
 amino acid
 Application US/09285310
 200154
 225 Franklin Street
 1410 amino acide
 617/542-8906
 100.0%; ilarity 100.0%; Conservative 0
 Lemontt, Jeffrey F.
CONCINO, Michael F.
VENTION: CHIMERIC PROTEINS FOR USE
VENTION: IN TRANSPORT OF A SELECTED SUBSTANCE INTO CELLS
 Conservative
 Heartlein, Michael W.
 unknown
 Fish & Richardson P.C.
 100.0%; Score 68; DB 2; I
100.0%; Pred. No. 0.0026;
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 US/09/285,310
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 07236/009002
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 Score 68; DB 2; : Pred. No. 0.0026;); Mismatches 0;
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 Length 1410;
 Indels
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 0
 Gaps
 Gaps
 0
 0
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밁

1378

YEKYLGEBYVKAV 13 ||||||||||||| YEKYLGEBYVKAV 1390

US-09-753-385-4

```
APPLICANT: Lemonit, Jeffrey F.

APPLICANT: Concino, Michael F.

TITLE OF INVENTION: CHIMERIC PROTEINS FOR USE IN TRANSPORT TITLE OF INVENTION: OF A SELECTED SUBSTANCE INTO CELLS

FILE REFERENCE: 10278-025004

CURRENT APPLICATION NUMBER: US/09/753,385

CURRENT FILING DATE: 2002-12-10

PRIOR APPLICATION NUMBER: US 08/470,058

PRIOR FILING DATE: 1995-06-06

PRIOR FILING DATE: 1995-06-06

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/037,188

PRIOR APPLICATION NUMBER: US 09/285,310
 밁
 S
 ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-753-385-4
 US-08-145-681-6
 RESULT 12
 Sequence 6, App. No. 557169
 SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 1410
 Sequence 4, Application US/09753385 Patent No. 6858578
 Matches
 Query Match
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PIBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/145,681
 GENERAL INFORMATION:
APPLICANT: Connee
 APPLICANT: Heartlein, Michael W. APPLICANT: Lemontt, Jeffrey F. APPLICANT: Concino, Michael F.
 NUMBER OF SEQ ID NOS:
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Baker & Botts, L.L.P.
STREET: 910 Louisiana St
 APPLICANT: Conneely, Orla M.
APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: May Gregory S.
TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using cDN
TITLE OF INVENTION: Various Organsims
 ATTORNEY/AGENT INFORMATION:
 NUMBER OF SEQUENCES:
 Local Similarity
nes 13; Conserva
 FILING DATE:
CLASSIFICATION: 435
 CITY: Houston
STATE: Texas
 COUNTRY:
 1378 YEKYLGEEYVKAV 1390
 INFORMATION:
 1 YEKYLGEEYVKAV 13
 77002
McGregor, Martin L.
 Application US/08145681
 USA
 Conservative
 100.0%; Score 68; DB 2;
100.0%; Pred. No. 0.0026;
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 Mismatches
 Version
 DB 2; Length 1410;
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 Gaps
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US-08-453-703-6
 US-08-145-681-6
 Sequence 6, Appendix Patent No. 57
 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/453,703
FILING DATE: Concurrently herewith
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,681
FILING DATE: October 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8206-024
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
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Best Local Similarity
 TELEFAX: 415-854-3694
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
 TELEFAX: 713/229/1522
INFORMATION FOR SEQ ID NO:
 APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Producti
TITLE OF INVENTION: And Lact
TITLE OF INVENTION: Various
NUMBER OF SEQUENCES: 14
 GENERAL INFORMATION:
 ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 APPLICANT: CONNECT, DENIS R. APPLICANT: Headon, Denis R. APPLICANT: O'Malley, Bert W.
 ORIGINAL SOURCE:
ORGANISM: Sus
 MOLECULE TYPE: protein HYPOTHETICAL: YES ANTI-SENSE: NO
 TELECOMMUNICATION INFORMATION: TELEPHONE: 713/229/1874
 MOLECULE TYPE: protein HYPOTHETICAL: YES
 CORRESPONDENCE ADDRESS:
 SEQUENCE CHARACTERISTICS:
 REGISTRATION NUMBER: 29,329
REFERENCE/DOCKET NUMBER: 19928-0125
 STREET: 1155 Avenue of the Americas CITY: New York
 TOPOLOGY:
 COUNTRY:
 TYPE: amino acid
 TELEPHONE: 415-854-3660
 ADDRESSEE:
 ropology:
 ENGTH:
 ENGTH:
 671 YEKYLGSEYVTAI 683
 10; Conservative
 1 УЕКУЬСЕЕУУКАУ 13
 amino acid
 Application US/08453703
 New York
 703 amino acids
 703 amino acids
 713/229/1522
 USA
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 Sus scrofa
 linear
 linear
 Pennie & Edmonds
 82.4%;
76.9%;
 Various Organisms
 Production of Recombinant Lactoferrin and Lactoferrin Polypeptides Using cDNA Sequences in
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US-08-453-703-6
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 US-08-456-106-6
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 703 amino acids
TYPE: amino acid
TOPOLOGY: linear
 Sequence 6, Myr-
Sequence 6, Myr-
No. 5849881
 GENERAL INFORMATION:
APPLICANT: Conneel
APPLICANT: Headon,
APPLICANT: O'Malle
APPLICANT: May, Gr
 Query Match
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 APPLICATION NUMBER: US/08/456,106
FILING DATE: Concurrently herewith
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,681
APPLICATION NUMBER: 08/145,681
FILING DATE: October 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8206-025
 ORIGINAL SOURCE:
ORGANISM: Sus
 MOLECULE TYPE: protein HYPOTHETICAL: YES ANTI-SENSE: NO
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
 APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Production of Recombinant Lactoferrin
TITLE OF INVENTION: and Lactoferrin Polypeptides Using cDNA Sequences
TITLE OF INVENTION: Various Organisms
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie &
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 NUMBER OF SEQUENCES:
 STATE: New York
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Local Similarity 76.9%;
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 Application US/08456106
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76.9%;
 Orla M.
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 Gaps
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 US-08-456-108-6
 US-08-456-108-6
 RESULT 15
 GENERAL INFORMATION:
APPLICANT: Conneely, Orla M.
APPLICANT: Headon, Denis R.
APPLICANT: O'Malley, Bert W.
APPLICANT: May, Gregory S.
TITLE OF INVENTION: Production
TITLE OF INVENTION: Various Or.
 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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CURRENT APPLICATION DATA:
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FILING DATE: Concurrently herewith
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APPLICATION NUMBER: 08/145,681
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ATTORNEY/AGENT INFORMATION:
NAME: Albert P. Halluin
DEGITEPATTON NUMBER: 25/27
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REFERENCE/DOCKET NUMBER: 821
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 CITY: New York
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 ADDRESSEE: Pennie & STREET: 1155 Avenue
 TYPE: amino acid
TOPOLOGY: linear
 REGISTRATION NUMBER:
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 671 YEKYLGSEYVTÁI 683
 1 УЕКУЦСЕЕУУКАУ 13
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 Production of Recombinant Lactoferrin
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 Listing first 45 summaries
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US-10-378-094-16
US-10-231-494-27
US-10-383-201-12
US-09-891-126-5
US-10-266-745-5
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 TYPE: PRT
 LENGTH:
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|------------------|-----------------|------------------|------------------|-------------------|-------------------|------------------|------------------|------------------|-------------------|-------------------|------------------|------------------|-----------------|------------------|------------------|------------------|------------------|
| 51               | 51              | 51               | 51               | 51                | 51                | 5<br>6           | 56               | 56               | 68                | 68                | 68               | 68               | 68              | 68               | 68               | 68               | 68               |
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| 708              | 708             | 708              | 698              | 698               | 695               | 703              | 15               | 15               | 1418              | 1410              | 1410             | 1074             | 869             | 698              | 698              | 698              | 698              |
| σ                | v               | 4.               | 4.               | 4.                | 4                 | ហ                | ഗ                | v                | σ                 | ຫ                 | w                | w                | 0               | 0                | v                | <sub>5</sub>     | 4                |
| US-10-513-523-2  | US-10-620-256-4 | US-10-169-297-8  | US-10-205-331-55 | US-10-316-253-275 | US-10-316-253-273 | US-10-620-256-6  | US-10-769-514-47 | US-10-769-514-14 | US-10-473-127-804 | US-10-473-127-805 | US-09-753-385-4  | US-09-753-385-2  | US-11-038-901-2 | US-11-177-506-46 | US-10-513-523-3  | US-10-887-711-2  | US-10-383-201-14 |
| Sequence 2, App. | Sequence 4, App | Sequence 8, App. | Sequence 55, App | Sequence 275, Ap  | Sequence 273, Ap  | Sequence 6, App. | Sequence 47, App | Sequence 14, App |                   | Sequence 805, App | Sequence 4, App. | Sequence 2, Appl | Sequence 2, App | Sequence 46, App | Sequence 3, Appl | Sequence 2, App. | Sequence 14, App |
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## ALIGNMENTS

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Sequence 4, Application US/10612162

| Publication No. US2004001415A1
| GENERAL INFORMATION:
| APPLICANT: Dade Behring Marburg GmbH
| TITLE OF INVENTION: antibodies, their preparation and use
| TITLE OF INVENTION: antibodies, their preparation and use
| CURRENT FILING DATE: 2002-07-03
| PRIOR APPLICATION NUMBER: US/10/612,162
| CURRENT FILING DATE: 2003-05-19
| RIOR FILING DATE: 2003-05-19
| NUMBER OF SEQ ID NOS: 4
| SOFTWARE: Patentin Ver. 2.1
| SEQ ID NO 4
| SEGUENCE OF SEQ ID NOS: 4
| SOFTWARE: PRIOR DATE: 2003-05-19
| NUMBER OF SEQ ID NOS: 4
| SOFTWARE: PRIOR DATE: 2003-05-19
| NUMBER OF SEQ ID NOS: 4
| SOFTWARE: PRIOR DATE: 2003-05-19
| NUMBER OF SEQ ID NOS: 4
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| TYPE: PRI
| OURLY MATCH | 100.0%; Score 68; DB 4; Length 13;
| Best Local Similarity | 100.0%; Fred. No. 0.0002; Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps
| USANISM: Homo sapiens
| US-10-378-094-16 | US/10378094 |
| PRIOR APPLICANT: PRIOR, Christopher P. APPLICANT: AND CONSERVENCY 13 |
| PRIOR MATCH | 100.0%; PRIOR DATE: 2003-03-04 |
| PRIOR APPLICANT: TURNER, Andrew P. APPLICANT: TURNER, Andrew DIFLED TRANSFERIN FUSION PROTEINS |
| TITLE OF INVENTION NUMBER: US/10/378,094 |
| CURRENT APPLICANTION NUMBER: US/10/378,094 |
| CURRENT FILING DATE: 2003-03-04 |
| PRIOR APPLICANTION NUMBER: US/0/314,494 |
| PRIOR APPLICANTION NUMBER: US/0/315,745
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; ORGANISM: Homo sapiens
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US-10-378-094-16
 APPLICANT: SADEGHI, Homayoun
APPLICANT: PRIOR, Christopher P.
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS
FILE REFERENCE: 54710-5004-US
CURRENT APPLICATION NUMBER: US/10/384,060
CURRENT FILLING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/334,759
PRIOR APPLICATION NUMBER: US 60/315,745
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PRIOR APPLICATION NUMBER: US 60/406,977
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PRIOR APPLICATION NUMBER: US 60/406,977
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 ; OTHER INFORMATION: C2 domain of huma US-10-384-060-38
 US-10-231-494-27
Sequence 27, Application US/10231494
Publication No. US20040023334A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
CURRENT APPLICATION NUMBER: US/10/231,494
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/315,745
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PRIOR APPLICATION NUMBER: US 60/334,059
 SOFTWARE: PatentIn version 3.2
SEQ ID NO 38
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TYPE: PRT
 Sequence 38, Application US/10384060 Publication No. US20030226155A1 GENERAL INFORMATION:
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 NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn version 3.2
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 Matches
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 PRIOR FILING DATE: 2001-08-30
 FEATURE:
 ORGANISM: Artificial sequence
 ENGTH: 49
 Local Similarity 100
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 37
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 1 YEKYLGEEYVKAV 13
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 Score 68;
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US-10-231-494-27
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 US-10-383-201-12
 CURRENT FILING DATE: 2003-03-06
PRIOR APPLICATION NUMBER: 10/029020
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/365,984
PRIOR FILING DATE: 2002-03-20
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PRIOR APPLICATION NUMBER: 60/389,143
PRIOR FILING DATE: 2002-06-14
PRIOR FILING DATE: 2002-06-14
 PRIOR APPLICATION NUMBER: 60/410,755
PRIOR FILING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: 60/412,957
PRIOR FILING DATE: 2002-09-23
PRIOR APPLICATION NUMBER: 10/051,874
PRIOR FILING DATE: 2002-01-16
PRIOR APPLICATION NUMBER: 60/366,928
PRIOR APPLICATION NUMBER: 60/366,928
 SOFTWARE: CuraSeqList version SEQ ID NO 12 LENGTH: 137
 PRIOR FILING DATE: 2001-
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PatentIn Ver.
SEQ ID NO 27
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 GENERAL INFORMATION:
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 FILE REFERENCE: 21402-568A CURRENT APPLICATION NUMBER: US/10/383,201 CURRENT FILING DATE: 2003-03-06
 PRIOR FILING DATE: 2002-03-22
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PRIOR FILING DATE: 2002-01-22
 NUMBER OF SEQ ID NOS:
 PRIOR APPLICATION NUMBER: 60/391,779
PRIOR FILING DATE: 2002-06-26
 APPLICANT: Alsobrook II, John et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHODS
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105 YEKYLGEEYVKAV 117
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 RESULT 8
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 RESULT 7
US-10-266-745-5
 ; ORGANISM: Homo sapiens
US-10-266-745-5
 Sequence 3, Application US/10378094 Publication No. US20030221201A1 GENERAL INFORMATION:
 Sequence 5, Application US/10266745 Publication No. US20030149256A1 GENERAL INFORMATION:
 Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 13; Conservative 0
 Query Match 100.0%; Score 68; DB 4; Length 328; Best Local Similarity 100.0%; Pred. No. 0.0063;
 SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 5
 Matches
 APPLICANT: PRIOR, Christopher P.
APPLICANT: LAI, Char Huei
APPLICANT: SADEGHI, Homayoun
APPLICANT: TURNER, Andrew
TITLE OF INVENTION: MODIFIED TRANSFERRIN FUSION PROTEINS
FILE REFERENCE: 54710-5001-01-US
CURRENT APPLICATION NUMBER: US/10/378,094
CURRENT FILING DATE: 2003-03-04
 FILE REFERENCE: PT035P1
CURRENT APPLICATION NUMBER: US/09/891,126
CURRENT FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR APPLICATION NUMBER: 60/171,595
PRIOR FILING DATE: 1999-12-23
 CURRENT APPLICATION NUMBER: US/10/266,745
CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US/09/891,126
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: PCT/US00/34769
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
 APPLICANT: Ruben et al. Transferrin Polynucleotides, Polypeptides, and Antibodies FILE REFERENCE: PT035P1
 NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn Ver. 2.0
 APPLICANT: Ruben et al.
TITLE OF INVENTION: Transferrin Polynucleotides, Polypeptides, and Antibodies
 NUMBER OF SEQ ID NOS: 7
 PRIOR APPLICATION NUMBER: 60/171,595 PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: US 10/231,494
 LENGTH: 328
TYPE: PRT
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 296 YEKYLGEEYVKÁV 308
 13;
 1 YEKYLGEEYVKAV 13
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 1999-12-23
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 Score 68; DB 3;
Pred. No. 0.0063;
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 RESULT 10
US-10-231-494-3
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; OTHER INFORMATION: Mature Transferrin Protein
US-10-384-060-3
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Sequence 3, Application US/10231494
Publication No. US20040023334A1
GENERAL INFORMATION:
APPLICANT: Prior, Christopher P.
TITLE OF INVENTION: Modified Transferrin Fusion Proteins
 Query Match
Best Local Similarity
Matches 13; Conserv
 PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 80
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
 Query Match
Best Local Similarity
Matches 13; Conserv
 SEQ ID NO 3
 GENERAL INFORMATION:
 Sequence 3, Application US/10384060 Publication No. US20030226155A1
 PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 66
NUMBER OF SEQ ID NOS: 66
SOFTWARE: Patentin version 3.2
 PRIOR APPLICATION NUMBER: US 10/231,494
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR FILING DATE: 2001-08-30
 APPLICANT: TURNER, Andrew TITLE OF INVENTION: MODIFIED TRANSFERRIN-ANTIBODY FUSION PROTEINS FILE REFERENCE: 54710-5004-US CURRENT APPLICATION NUMBER: US/10/384,060 CURRENT FILING DATE: 2003-03-10
 APPLICANT: SADEGHI, Homayoun
APPLICANT: PRIOR, Christopher P.
APPLICANT: TURNER, Andrew
 LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
 TYPE: PRT
ORGANISM: Homo sapiens
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 LENGTH: 679
 647
 647 YEKYLGEEYVKAV 659
 1 УЕКУІСЕЕУУКАУ 13
 1 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 659
 Conservative
 Conservative
 100.0%;
 100.0%;
 <u>,</u>
 0
 Score 68; DB 4; Pred. No. 0.014;
 Score 68; DB 4;
Pred. No. 0.014;
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 DB 4;
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 Length 679;
 Length 679;
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FILE REFERENCE: 54710-5001-US

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CURRENT APPLICATION NUMBER: US/10/231,494
CURRENT FILING DATE: 2002-08-30
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PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: US 60/334,059
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 30
SOFTMARE: PATENTIN Ver. 2.1
SEQ ID NO 3
LENGTH: 679
 Sequence 4, Application US/10429482
Publication No. US20040219097A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The Diagnosis, Imaging and Treatment Of Tu
FILE REFERENCE: 2537.000001
CURRENT APPLICATION UMBER: US/10/429,482
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
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US-10-429-497-4
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 US-10-429-482-4
 ; ORGANISM: Homo sapiens
US-10-429-497-4
 US-10-231-494-3
Best Loc
Matches
 APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Methods For The Treatment of Tumors
FILE REFERENCE: 2537.000006
CURRENT APPLICATION NUMBER: US/10/429,497
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1
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LENGTH: 679
TYPE: PRT
 Sequence 4, Application US/10429497 Publication No. US20040219098A1 GENERAL INFORMATION:
 Query Match 100.0%; Score 68; DB 5; Length 679 Best Local Similarity 100.0%; Pred. No. 0.014;
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Best Local Similarity
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Best Local &
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 OTHER INFORMATION: Mature transferrin protein
 FEATURE:
 ORGANISM: Homo sapiens
 TYPE: PRT
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Sequence 4, Application US/10429515

Publication No. US20040219099A1

GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
ITILE OF INVENTION: Methods For The Treatment Of Tumors
FILE REFERENCE: 2537.000005

CURRENT APPLICATION NUMBER: US/10/429,515

CURRENT FILING DATE: 2003-05-02

NUMBER OF SEQ ID NOS: 6

SOFTWARE: Patentin version 3.1

SEQ ID NO 4
 Sequence 4, Application US/10429598
Publication No. US20040219100A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For The FILE REFERENCE: 2537.000003
CURRENT APPLICATION NUMBER: US/10/429,598
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
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 RESULT 14
 US-10-429-515-4
 US-10-429-515-4
 RESULT 15
Sequence 4, Application US/10429635
Publication No. US20040219101A1
GENERAL INFORMATION:
APPLICANT: Sandhu, Jasbir
TITLE OF INVENTION: Composition Useful For Treatment Of Tumors
FILE REFERENCE: 2537.000007
CUCRENT APPLICATION NUMBER: US/10/429,635
CURRENT FILING DATE: 2003-05-02
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
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TYPE: PRT
ORGANISM: Homo sapiens
 LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No.
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Pred. No. 0.014;
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US-11-258-767-15

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| g Qy                                                  | Query Matc<br>Best Local<br>Matches         | ; OTHER IN<br>; OTHER IN<br>; OTHER IN<br>US-10-515-42 | LENGTH: 4 ; TYPE: PRT ; ORGANISM: | TWAI S | ; PRIOR I               | ; PRIOR I                |                            | ; PRIOR !                               | 22         | ; FILE RI      | ; APPLICANT:                            | GENERAL INF | RESULT 1 US-10-515-4 ; Sequence ; highlight |            | 4.4          |            | 41          | 4 S<br>0 9 | 38       | 36         | ນ ພ<br>4 7 | ω i<br>ω i | 331                      | 30         | 28      | 26<br>27               |  |
|-------------------------------------------------------|---------------------------------------------|--------------------------------------------------------|-----------------------------------|--------|-------------------------|--------------------------|----------------------------|-----------------------------------------|------------|----------------|-----------------------------------------|-------------|---------------------------------------------|------------|--------------|------------|-------------|------------|----------|------------|------------|------------|--------------------------|------------|---------|------------------------|--|
| 1 YEKYLG<br>      <br>37 YEKYLG                       | h<br>Sim<br>13;                             | INFORMATION:<br>INFORMATION:<br>109-27                 | 9<br>Artifi                       | Patent | JA IO                   | APPLICATION FILING DATE: |                            | APPLICATION NUMBER: FILING DATE: 2003-0 |            | REFERENCE: 054 | NT: Turner, I                           | NFORMAT     | 29-27<br>27, Api                            |            | თ თ          | , 7 .      | 1 7         | 5 U        | 58.      | 5 66.      | 5 66.      | 7 69.      | 8 70.<br>8 70.           | 70.        | 8 70.   | 70.                    |  |
| YEKYLGEEYVKAV 13<br>             <br>YEKYLGEEYVKAV 49 | 100.<br>ilarity 100.<br>Conservative        | N: Descript<br>N: peptide                              | cial Seque                        | Ver    | NUMBER: U<br>: 2003-07- |                          | NUMBER: US<br>: 2002-08-30 | œ                                       | <b>K</b> 4 | 054710-5007-WO | •                                       | hristo      | ion US                                      |            | -            | 618        |             | 94<br>94   | 414      | 707        |            | 696        | 711<br>711               | 711        | 711     | 711<br>711             |  |
|                                                       | 0%; Score<br>0%; Pred.<br>0; Mis            | tion of Art<br>insertion                               | nce                               |        | 60/485,<br>9            | 10/384,0                 | 60/406,<br>0               | PCT/US03/26779                          | /10/       | -WO            | ; = .                                   | er P.       | 3/10515429<br>187A1                         | ALIG       | US-10-4      | US-10-4    | US-10-      | 10-        | US-10-   | US-11-     | US-10-     | US-10-     | US-11-                   | US-11-     | US-11-2 | US-11-2                |  |
|                                                       | ore 68; DB 6<br>ed. No. 1.9e-<br>Mismatches | tificial<br>region                                     |                                   |        | 404                     | 60                       | 977                        | 779                                     | ,429       | FIOCETH        | d<br>1<br>2<br>3                        |             |                                             | ALIGNMENTS | 71-571A-4786 | 1-571A-    | 370-959-111 | 429-6      | 571A     | 767-       | 429-       | 129-       | 767-<br>767-             | 767-       | 8-767-1 | 58-767-15<br>58-767-17 |  |
|                                                       | ; Length<br>05;<br>0; Indel                 | Sequence:                                              |                                   |        |                         |                          |                            |                                         |            | PIDIGITES      | היא היא היא היא היא היא היא היא היא היא |             |                                             |            |              |            |             |            | 0        |            |            |            |                          |            |         |                        |  |
|                                                       | 49;                                         | Transferrin                                            |                                   |        |                         |                          |                            |                                         |            |                |                                         |             |                                             |            |              | Sequence 4 | Sequence 1  | Sequence ( | Sequence | Sequence 1 | Sequence 3 | Sequence 4 | Sequence 3<br>Sequence 3 | Sequence 2 | equence | Sequence 1             |  |
|                                                       | Gaps 0;                                     | ន៍                                                     |                                   |        |                         |                          |                            |                                         |            |                |                                         |             |                                             |            | 4786, Ap     | A          | -           | 67, Appl   | 6<br>A   |            | App        |            | App                      |            | App     | 5, Appl                |  |

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US-10-515-429-3
; Sequence 3, Application US/10515429
; Publication No. US20060105387A1
; GENERAL INFORMATION:
; APPLICANT: Prior, Christopher P.
; APPLICANT: Turner, Andrew J.
; APPLICANT: Sadeghi, Homayoun

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APPLICANT: PTIOR, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-08-30
PRIOR FILING DATE: 2003-03-10
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER: US 60/485,404
 FITTLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR FILING DATE: 2003-08-30
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
NUMBER OF SEQ ID NOS: 75
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US-10-515-429-37
; Sequence 37, Application US/10515429
; Publication No. US20060105387A1
; GENERAL INFORMATION:
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 ; FEATURE:
; OTHER INFORMATION: Mature transferrin protein
US-10-515-429-3
 ; ORGANISM: Homo sapiens US-10-515-429-2
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 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 698
TYPE: PRT
 sequence 2, Application US/10515429
Publication No. US20060105387A1
GENERAL INFORMATION:
APPLICAME S.
 Query Match 100.0%; Score 68; Best Local Similarity 100.0%; Pred. No. Matches 13; Conservative 0; Mismatch
 SEQ ID NO 3
LENGTH: 679
TYPE: PRT
ORGANISM: Homo sapiens
 Matches
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Best Local Similarity
 SOFTWARE: PatentIn Ver. 2.1
 NUMBER OF SEQ ID NOS: 75
 666 YEKYLGEEYVKAV 678
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; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 36
; LENGTH: 686
; TYPE: PRT
; ORGANISM: PIG
US-11-258-767-36
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 ; TYPE: PRT ; ORGANISM: Oryctolagus cuniculus US-10-515-429-37
 US-11-258-767-37
 US-11-258-767-36
 PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR PPLICATION NUMBER: US 60/406,977
PRIOR PILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 60/485,404
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
 Sequence 36, Application US/11258767

Publication No. US20060094082A1

GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Varadhachary, Atul
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67

NUMBER OF SEQ ID NOS: 67
Sequence 37, Application US/11258767 Publication No. US20060094082A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn Ver.
SEQ ID NO 37
LENGTH: 676
 Matches
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 Query Match
 APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-W0
CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11.23
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 85.3%;
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 Score 56; DB 7;
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 Gaps
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APPLICANT: Varadhachary, Atul APPLICANT: Glynn, Peter APPLICANT: Petrak, Karel

Glynn, Peter Petrak, Karel Engelmayer, Jose

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 ; TYPE: PRT
; ORGANISM: PIG
US-11-258-767-37
 US-11-258-767-32
 ORGANISM: GOAT
 Sequence 32, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Varadhachary, Atul
APPLICANT: Engelmayer, Jose
APPLICANT: Engelmayer, Jose
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
TILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER: TILING DATE: 2004-10-26
 NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
SEQ ID NO 28
LENGTH: 708
TYPE: PRT
 Sequence 28, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 37 LENGTH: 703
 Matches
 Query Match
 APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR APPLICATION NUMBER: US 60/622,176
 Query Match
 PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
 APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
 PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
 CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
 FILE REFERENCE: HO-P02915US1
 TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
 Local
 Local
 671 YEKYLGSEYVTAI 683
 676 YEKYLGTEYVTAI 688
 10;
 10;
 1 УЕКУЬСЕЕУУКАУ 13
 1 YEKYLGEEYVKAV 13
 Similarity
 Similarity
 Conservative
 Conservative
 80.9%;
 82.4%;
 Score 55; DB 7;
Pred. No. 0.051;
1; Mismatches
 Score 56; DB 7;
Pred. No. 0.034;
 DB 7;
 DB 7;
 Length 708;
 2;
 Length 703;
 ٥,
 0
 Gaps
 Gaps
 0
 0
 á
 밁
 ; TYPE: PRT
; ORGANISM: GOAT
US-11-258-767-32
 US-11-258-767-13
 맑
 ; ORGANISM: Rattus norvegicus US-10-515-429-38
 CURRENT APPLICATION NUMBER: US/10/515,429
CURRENT FILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: PCT/US03/26779
PRIOR FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US 60/406,977
PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR APPLICATION NUMBER: US 10/384,060
PRIOR FILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-03-10
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 75
LENGTH: 681
TYPE: PRT
ORGANISM: COW
IS-11-258-767-13
 GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 13
 Sequence 13, Application US/11258767 Publication No. US20060094082A1
 Matches
 SEQ ID NO 38
 Query Match
 GENERAL INFORMATION
 Sequence 38, Ap
Publication No.
 Matches
 APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
 Best Local Similarity
Matches 9; Conserv
 Query Match
Best Local Similarity
 APPLICANT: Prior, Christopher P.
APPLICANT: Turner, Andrew J.
APPLICANT: Sadeghi, Homayoun
TITLE OF INVENTION: Transferrin Fusion Protein Libraries
FILE REFERENCE: 054710-5007-WO
 LENGTH: 67
 SOFTWARE:
 LENGTH: 708
 644
 676 YEKYLGTEYVTÁI 688
 10;
 1 YEKYLGEEYVKAV 13
 1 YEKYLGEEYVKAV 13
 PatentIn Ver. 2.1
 YEEYLGAEYLQAV 656
 Application US/10515429
 Conservative
 Conservative
 US20060105387A1
 75.0%;
69.2%;
 80.9%;

 Mismatches

 Pred.
 Score 51; I
Pred. No. 0.
 Score 55; DB 7;
Pred. No. 0.051;
 Mismatches
 DB 6;
 DB 7;
 Length 676;
 Length 708;
 Indels
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 Gaps
 Gaps
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0,

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NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
SEQ ID NO 25
LENGTH: 698
TYPE: PRT
ORGANISM: RAT
US-11-258-767-25
В
 RESULT 12
US-11-258-767-16
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 RESULT 11
US-11-258-767-25
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 ORGANISM: COW
US-11-258-767-16
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 16
LENCTH: 708
TYPE: PRT
 Sequence 16, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:
 Query Match
Best Local :
 Sequence 25, Application US/11258767
Publication No. US20060094082A1
 Query Match
Best Local Similarity
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Petrak, Karel
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITILE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
 APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR PILIAGION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
 APPLICANT:
 APPLICANT: Varadhachary, Atul
 Local
 649
 666 YEEYLGAEYLQAV 678
 1 YEKYLGEEYVKAV 13
 1 YEKYLGEEYVKAV 13
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 9;
 Similarity 69.: 9; Conservative
 YEKYLGEEYVKAV 13
 YEEYLGTEYVTAI 661
 YEEYLGTEYVTAI 688
 Glynn, Peter
Petrak, Karel
 75.0%;
ilarity 69.2%;
Conservative
 Conservative
 75.0%;
69.2%;
 75.0%;
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 Score 51; DB
Pred. No. 0.23
2; Mismatches
 Score 51; DB
Pred. No. 0.24
3; Mismatches
 Score 51; DB 7;
Pred. No. 0.24;
2; Mismatches
 DB
1.24;
 DB 7;
 ۲,
 2; Indels
 Length 708;
 Length 681;
 Length 698;
 Indels
 Indels
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 Gaps
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 Query Match
Best Local Similarity
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 Query Match
Best Local Similarity
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 ; ORGANISM: COW
US-11-258-767-22
 US-11-258-767-29
 RESULT 15
 US-11-258-767-22
 RESULT 14
 US-11-258-767-19
 US-11-258-767-19
 Sequence 19, Application US/11258767

Publication No. US20060094082A1

GENERAL INFORMATION:

APPLICANT: Varadhachary, Atul

APPLICANT: Glynn, Peter

APPLICANT: Petrak, Karel

APPLICANT: Petrak, Karel

APPLICANT: Engelmayer, Jose

TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF

FILE REFERENCE: HO-P02915US1
 Sequence 22, Application US/11258767
Publication No. US20060094082A1
GENERAL INFORMATION:
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
 CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.3
SEQ ID NO 19
 Sequence 29, Application US/11258767
Publication No. US20060094082A1
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 22
LENGTH: 708
 GENERAL INFORMATION:
 APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
APPLICANT: Engelmayer, Jose
TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
FILE REFERENCE: HO-P02915US1
CURRENT APPLICATION NUMBER: US/11/258,767
CURRENT FILING DATE: 2005-10-26
PRIOR APPLICATION NUMBER: US 60/622,176
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
APPLICANT: Varadhachary, Atul
APPLICANT: Glynn, Peter
APPLICANT: Petrak, Karel
 NUMBER OF SEQ ID NOS: 67
 LENGTH: 708
TYPE: PRT
ORGANISM: WATER BUFFALO
 TYPE: PRT
 676 YEEYLGTEYVTAI 688
 676 YEEYLGTEYVTAI 688
 1 YEKYLGEEYVKAV 13
 YEKYLGEEYVKAV 13
 Conservative
 Conservative
 75.0%;
69.2%;
 75.0%;
 2
 Score 51; DB 7;
Pred. No. 0.24;
 Score 51; I
Pred. No. 0.
 Mismatches
 Mismatches
 DB 7;
 2
 Length 708;
 Length 708;
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 0,
 Gaps
 Gaps
 0
 0
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```
; APPLICANT: Engelmayer, Jose
; TITLE OF INVENTION: COMPOSITIONS OF LACTOFERRIN RELATED PEPTIDES AND USES THEREOF
; FILLE REFERENCE: HO-PO2915US1
; CURRENT APPLICATION UNMBER: US/11/258,767
; CURRENT FILING DATE: 2005-10-26
; PRIOR APPLICATION NUMBER: US 60/622,176
; PRIOR FILING DATE: 2004-10-26
; NUMBER OF SEQ ID NOS: 67
; SOFTWARE: Patentin version 3.3
; SEQ ID NO 29
; LENGTH: 708
; TYPE: PRT
; ORGANISM: COW
US-11-258-767-29

Query Match
Best Local Similarity 69.2%; Pred. No. 0.24;
Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

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Search completed: June 13, 2006, 04:56:52

Job time: 4.39706 secs
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Result
 Minimum
Maximum
 Run
 Post-processing: Minimum Match
Maximum Match
Listing first
 OM protein -
 Database
 Scoring table:
 Sequence:
 Perfect score:
 Title:
 9.
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 number of hits satisfying
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length:
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1: geneseqp198
2: geneseqp199
 Query
Match
 9:
10:
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 US-10-612-162A-4
68
 June 13, 2006, 03:02:32 ; Search time 194 Seconds
 2589679 seqs, 457216429 residues
 Gapop 10.0 , Gapext 0.5
 YEKYLGEEYVKAV 13
 Copyright
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geneseqp2003bs:*
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 geneseqp2006s:*
 Length
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 DВ
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100%
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AEE30609
AEF52225
 AAW70384
AEA79013
ADW33519
 ADP83124
ADU07237
 ADW34752
AAW21308
AAW21301
AAW21313
 ADT88295
AED66887
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ADX26787
 ADW39569
 ADW39581
 summaries
 SUMMARIES
 model
 (without alignments)
30.638 Million cell updates/sec
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 Adg46078 Human CDT
Adu07228 Human IL-
Aed66887 Human IL-
Aed66887 Human IL-
Aed66859 Human IL-
Aed66859 Human IL-
Aed66859 Human IL-
Aed70313 Bovine Se
Adw33519 HLA bindi
Adw33519 HLA bindi
Adw34752 HLA bindi
Aaw21301 Glucagon
Aaw21301 Glucagon
Aaw21301 Glucagon
Aaw21313 Glucagon
Adw39131 Glucagon
Adw39131 Glucagon
Adw39111 Glucagon
Adw39111 Glucagon
Adw39111 Glucagon
Adw39111 Human IL-
Adw39561 HWG COA r
Adw39576 HWG COA r
Adw39576 HWG COA r
Adw39576 HWG COA r
Adw39576 HWG COA r
 Description
 Aed66896 Human IL-
Represent
Interfaci
Rat cytos
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Human CDT peptide fragment

ADG46078; 25-MAR-2004

(first entry)

ADG46078 standard; peptide; 13

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| 45                 | 44                 | 43                 | 42                 | 41                | 40                | 39                | 38                | 37                | 36                | 35                | 34                 | 33                 | 32                 | 31                 | 30                 | 29                 | 28                 | 27                 | 26                 | 25                 | 24                 |
|--------------------|--------------------|--------------------|--------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
| 23                 | 23                 | 23                 | 23                 | 23                | 23                | 23                | 23                | 23                | 23                | 23                | 23                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 |
| ü                  | ω                  |                    | ü                  | 'n                |                   | Ψ                 | ü                 | 33.8              |                   |                   | •                  |                    | 35.3               | 35.3               |                    | 35.3               |                    | 35.3               | 35.3               |                    | 35.3               |
| 13                 | 13                 | 13                 | 13                 | 13                | 13                | 13                | 13                | 13                | 13                | 13                | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 | 13                 |
| N                  | N                  | N                  | N                  | N                 | N                 | N                 | N                 | N                 | N                 | N                 | _                  | 5                  | 10                 | Ģ                  | φ                  | φ                  | 9                  | 9                  | 7                  | Ų                  | ω                  |
| AAY10774           | AAW61555           | AAR98417           | AAR90274           | AAW21307          | AAW21305          | AAW21312          | AAW21309          | AAW21298          | AAW21302          | AAW21300          | AAP82510           | AEG03146           | AEE39361           | AEA44892           | AEA53535           | AEA53802           | ADZ37854           | ADY64440           | ADM08910           | ABJ10898           | AAB39978           |
| Aay10774 Peptide u | Aaw61555 Biotinyla | Aar98417 Interleuk | Aar90274 Ion-chann | Aaw21307 Glucagon | Aaw21305 Glucagon | Aaw21312 Glucagon | Aaw21309 Glucagon | Aaw21298 Glucagon | Aaw21302 Glucagon | Aaw21300 Glucagon | Aap82510 Viral inh | Aeg03146 CD200-spe | Aee39361 Human pro | Aea44892 Apolipopr | Aea53535 Novel hum | Aea53802 Novel hum | Adz37854 Human kin | Ady64440 Human cyt | Adm08910 Canine im | Abj10898 K+beta M6 | Aab39978 Anti-hIL1 |

## ALIGNMENTS

```
KW antibody; carbohydrate-deficient transferrin; CDT;
KW non-glycosylated transferrin; alcoholism; alcoholic; diagnosis; human.
XX
OS Homo sapiens.
XX
EP1378521-A1.
XX
PN EP1378521-A1.
XX
PD 07-JAN-2004.
XX
PD 07-JAN-2004.
XX
PD 07-JAN-2004.
XX
PD 05-JUL-2002; 2003EP-00011334.
XX
PR 05-JUL-2002; 2002DE-01030550.
XX
PA (DADE-) DADE BEHRING MARBURG GMBH.
XX
PA Althaus H;
```

This invention describes a novel antibody that, in aqueous solution, binds selectively to carbohydrate-deficient transferrin (CDT) without having to bind CDT to a solid phase. The invention also describes an antibody that binds selectively to CDT at regions containing the sequences VVARSWGGKEDLIVELL, TTEDSIAKINNOSADAMSLDGGF, SKLSWGSGLNLSERN and YEKYLGERYVKAV (ADG46075-ADG46078) and an immunoassay for detecting CDT using the antibodies of the invention. The antibodies of the invention are monoclonal, specifically from hybridomas DSM ACC2540 or 2541. The antibodies are obtained by immunisation with non-glycosylated transferrin, or a fragment, then generation of hybridomas by standard fusion of spleen and myeloma cells. Hybridomas are then selected for

New antibody specific for carbohydrate-deficient transferrin, useful for diagnosis of alcoholism, can bind its target in solution, eliminating need for immobilization.

WPI; 2004-073743/08.

Claim 4; SEQ ID NO 4; 21pp; German.

3333333<del>3</del>3

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Matches
 Query Match
Best Local
 The invention relates to a method of managing, treating or ameliorating a respiratory infection or its symptom in a human subject by administering an interleukin-9 (II-9) antagonist. The method further comprises administering to the subject an anti-respiratory syncytial virus (anti-RSV) antigen antibody. The anti-RSV antigen antibody is palivizumab. The method further comprises administering a leukotriene modifier, which is montelukast, zafirlukast, pranlukast or zileuton. Also disclosed are: a method of preventing the development, onset or progression of asthma or one or more asthma-like symptoms in a child, a method of preventing, managing, treating or ameliorating wheezing in a pre-term infant, infant or child, a method of preventing, managing, treating or ameliorating wheezing in a human subject, and a method of preventing, managing, treating or ameliorating wheezing in a human subject, and a method of preventing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, managing, mana
 Managing, infection syncytial
 production of antibodies that specifically bind CDT from the aqueous phase. The antibodies are useful for diagnosis of alcoholism, CDT, whic lacks glycosylation at AssAll and/or Asn 611, is present only in alcoholics. The antibodies allow direct detection of CDT in solution, eliminating the need for immobilising it on a solid phase (as required when using known antibodies). ADG46075-ADG46094 represent CDT peptide fragments used in the method of the invention.
 Respiratory infection; human; interleukin-9 antagonist; IL-9; respiratory syncytial virus; anti-RV antigen antibody; leukotriene modifier; asthma-like symptom; wheezing; allergy; immunomodulatory; anti-Inflammatory; anti-viral; antibiotic; antifungal; mast cell modulator; virucide; antibacterial; fungicide; antiasthmatic; antiallergic; 4D4 antibody; variable heavy chain; VH;
 Disclosure; SEQ ID NO 3; 276pp; English.
 WPI; 2004-775524/76.
 12-APR-2004; 2004WO-US011329
 28-OCT-2004.
 ADU07228 standard; peptide; 13 AA.
 Sequence 13 AA;
 11-APR-2003; 2003US-0462307P.
10-JUN-2003; 2003US-0477801P.
 Homo sapiens
 complementarity determining region; CDR
 Human 4D4 variable heavy chain (VH) CDR3.
 13-JAN-2005
 (MEDI-) MEDIMMUNE INC
 13;
 1 YEKYLGEEYVKAV 13
 1 YEKYLGEEYVKAV 13
 Similarity
 treating or ameliorating in a human subject a respiratory or its symptoms caused by bacteria, fungi or respiratory virus by administering an IL-9 antagonist.
 Conservative
 (first entry)
 100.0%; Score 68; Di
100.0%; Pred. No. 0.1
tive 0; Mismatches
 DB 8;
0.00047;
 0; Indels
 Length 13;
 0
 CDT, which
 Gaps
 0
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RESULT 3
ADT88295
 CC subject. Preventing the development, onset or progression of asthma or come or more asthma-like symptoms in a child having or previously having a Crespiratory infection comprises administering an IL-9 antagonist. CC Preventing, managing, treating or ameliorating wheezing in a pre-term CC infant, infant or child comprises administering an IL-9 antagonist. The cmethod further comprises administering at least one other therapy that is comethod further comprises administering at least one other therapy that is comediated in IL-9 antagonist. Preventing, managing, treating CC or ameliorating wheezing in a human subject comprises administering an IL-9 antagonist and at least one other therapy that is not administering an IL-9 antagonist. The therapy is an immunomodulatory agent, an anti-viral agent, an anti-blotic, an anti-fungal cadministering to the subject a leukotriene modifier, an anti-histody or a mast cell modulator. The method further comprises cadministering an IL-9 antagonist and a leukotriene modifier, an anti-histody or a mast cell protease inhibitor. Preventing, managing, treating or ameliorating cadministering an IL-9 antagonist is an anti-body, an anti-IL-4 antibody or a mast cell protease inhibitor. Preventing, managing, treating or ameliorating cadministering an IL-9 antagonist is an antibody that immunospecifically binds to an IL-9 antagonist is an antibody that immunospecifically binds to an IL-9 receptor (IL-9R) or its symptoms caused by bacteria, fungical competities and thousand the sequence represents a complementarity determining region con RSV. This sequence represents a complementarity determining region
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 S
 Query Match
Best Local
 Matches
 respiratory; antiinflammatory; immunosuppressive; antiasthmatic; antiallergic; antiarthritic; neuroprotective; antirheumatic; immunotoxin; angiogenesis inhibitor; TWF alpha antagonist; IL-9 antagonist; IL-9; IL-9; antibody; 4D4; 4D4 H2-1 D11; 4D4com-XF9; 4D4com-2F9; 7F3; 71A10; 22D3; 7F3com-2H2; 7F3com-3H5; 7F3com-3D4; IL-9 associated disorder; respiratory infection; respiratory syncytial virus; parainfluenza virus; human metapneumovirus; inflammatory disorder; asthma; allergy; arthritis; autoimmune disorder; rheumatoid arthritis; multiple sclerosis; autoimmune disorder; rheumatoid arthritis; multiple sclerosis;
 WPI; 2004-775520/76
 11-APR-2003; 2003US-0462259P.
10-JUN-2003; 2003US-0477797P.
 12-APR-2004; 2004WO-US011172
 Human IL-19 antibody variable heavy domain CDR3 seqid 3.
 13-JAN-2005
 ADT88295 standard; peptide; 13 AA
 Sequence 13 AA;
 WO2004091510-A2
 Homo sapiens
 variable heavy
 (MEDI-) MEDIMMUNE INC
 l Similarity
5; Conserv
 ω
 4 YLGEEYVK 11
 YYGSDYVK 10
 Conservative
 (first entry)
 domain; complementarity
 44.1%;
 Score 30;
Pred. No.
 DB 8; Le.. 6.1e+02;
 determining
 Length 13;
 0
 Gaps
 0
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New IL-9 antibody comprising a variable heavy or light domain

or

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RESULT 4
AED66807
ID AED6
XX AED6
XX AED6
XX AED6
XX AED6
XX Gene
XW Gene
XW anti
XW anti
XW anti
XW anti
XW anti
XW anti
XW AED6
XW Tesp
XW Fes
XW Fes
XW Heav
XW Tesp
XW Homc
XX Tesp
XX US20
XX US20
AED60
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 CC (VH) or variable light (VL) domain having an amino acid sequence of the CVH or VL domain of 4D4, 4D4 H2-1 D11, 4D4com-XF9, 4D4com-ZF9, 7F3, 71A10, CC 2D3, 7F3com-2H2, 7F3com-3H5 or 7F3com-3D4; or a complementarity CC determining region (CDR) having an amino acid sequence of a CDR of 4D4, 4D4 H2-1 D11, 4D4com-XF9, 4D4com-XF9, 7F3, 71A10, 2D3, 7F3com-3H5 or 7F3com-3D4. The antibody immunospecifically binds to a CC comprising the II-9 antibody and a carrier; a method for preventing, cc treating, managing or ameliorating a disease or disorder characterised by aberrant expression or activity of an II-9 polypeptide or an II-9R; a cc method of diagnosing, prognosing, or monitoring a disorder or disease characterised by aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising and an activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising and an activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC or activity of an II-9R; a composition for diagnosing, treating or composition for diagnosing, treating or composition for diagnosing, treating or caused by bacteria or respiratory syncytial virus, parainfluenza virus or human metappeneumovirus, inflammatory disorder such as asthma, allergy or antibody variable heavy domain composition acid sequence of a human II-19
 Query Match
 Gene therapy; chemotherapy; animal disease model; inflammation; antiinflammatory; asthma; antiasthmatic; allergy; antiallergic; autoimmune disease; immunosuppressive; diabetes; antidiabetic; hashimotos disease; antithyroid; adrenal disease; endocrine-gen.; anem. antianemic; multiple sclerosis; neuroprotective; heumatoid arthritis; antiarthritic; antirheumatic; systemic lupus erythematosus; dermatological; carditis; cardiant; sjoegrens syndrome; polymyositis; muscular-gen.; dermatomyositis; scleroderma; hyperproliferation; cytostatic; infection; antimicrobial; respiratory tract infection; respiratory-gen.; interleukin-9; cytokine; lymphokine; antibody; heavy chain variable region; 404 H2-1 D11; 7F3; 71A10; 7F3 22D3;
 complementarity determining region having an amino acid sequence of e.g., 4D4, useful in treating or preventing e.g., respiratory infection or inflammatory disorder.
 Human IL-9 antibody,
 Sequence 13
 Disclosure; SEQ ID NO 3; 291pp; English.
 26-JAN-2006
 AED66887 standard; peptide; 13 AA
 antibody variable heavy domain complementarity determining region
 Local
 invention describes a new IL-9 antibody comprising: a variable heavy) or variable light (VL) domain having an amino acid sequence of the or VL domain of 4D4, 4D4 H2-1 D11, 4D4com-XF9, 4D4com-2F9, 7F3, 71A10
 4 YLGEEYVK 11
 44.1%;
Similarity 62.5%;
5; Conservation
 YYGSDYVK 10
 3
 (first entry)
 4D4 H2-1 D11 VH domain CDR3 peptide,
 Score 30; DB 8; Le
Pred. No. 6.1e+02;
1; Mismatches 2;
 Length 13;
 SEQ ID:
 0,
 Gaps
 anemia;
 0
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24-NOV-2005

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RESULT 5
AED66959
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 CC polypepide, IL-9R or one or more subunits, inflammatory disorders such as asthma or an allergy, autoimmune disorders such as diabetes, autoimmune adrenal insufficiency, pure red cell anemia, multiple sclerosis, rheumatoid carditis, systemic lupus crythematosus, rheumatoid arthritis, chronic inflammation, Sjogren's syndrome polymyositis, dermatomyositis, chronic inflammation, Sjogren's syndrome polymyositis, dermatomyositis, chronic inflammation, Sjogren's cyndrome polymyositis, dermatomyositis, a respiratory infection). The disease or an infection (preferably, a respiratory infection). The crimention is also useful in gene therapy, chemotherapy and to generate animal disease models. The present sequence is a heavy chain variable (VH) domain complementarity determining region, CDR3 peptide present in human interleukin-9 (IL-9) antibodies 4D4 H2-1 D11, 7F3, 71A10, 7F3 2D3, 7F3com-3H2, 7F3com-3H5 and 7F3com-3D4. Note: The present sequence is the SEQ ID NO: 3 which is given in the sequence listing. This sequence C differs from the SEQ ID NO: 3 shown on table 1 in page 2 (antibody 4D4) circle in the disclosure of the specification (see AED66959).
 Matches
 Query Match
Best Local
Gene therapy; chemotherapy; animal disease model; inflammation; antiinflammatory; asthma; antiasthmatic; allergy; antiallergic; autoimmune disease; immunosuppressive; diabetes; antidiabetic; hashimotos disease; antithyroid; adrenal disease; endocrine-gen.; anemi antianemic; multiple sclerosis; neuroprotective; rheumatoid arthritis; antirheumatic; systemic lupus erythematosus; dermatological; carditis; cardiant; sjoegrens syndrome; polymyositis; muscular-gen.; dermatomyositis; scleroderma; hyperproliferation; cytostatic; infection; antimicrobial; respiratory tract infection;
 The present invention provides liquid formulation of antibodies or antibody fragments that immunospecifically binds to an interleukin-9 (II-9) polypeptide. The formulation exhibit stability, low to undetectable levels of aggregation and very little to no loss of the biological activities of antibodies and its fragments, even during the long period of storage. The antibody formulation of the invention is useful for preventing, managing, treating or ameliorating a disease or disorder associated with aberrant expression and/or activity of an II-9
 AED66959 standard; peptide; 13
 New antibody formulation comprises antibody or antibody fragment that immunospecifically binds to an IL-9 polypeptide, useful for preventing, managing, treating, or ameliorating, e.g. asthma, allergy, diabetes, or multiple sclerosis.
 WPI; 2005-796072/81.
N-PSDB; AED66930.
 Sequence 13
 Disclosure;
 Human IL-9 antibody, 4D4 VH domain CDR3 peptide,
 26-JAN-2006 (first entry)
 12-APR-2004; 2004US-0561845P
 12-APR-2005; 2005US-00105268
 (MEDI-) MEDIMMUNE INC
 ω
 4 YLGEEYVK 11
 ა
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 Similarity
 YYGSDYVK 10
 Conservative
 A,
 SEQ ID NO
 44.1%;
62.5%;
 3; 125pp; English.
 Score 30; DB 9;
Pred. No. 6.1e+02;
 Mismatches
 N
 Length 13
 SEQ ID:
 0
 Gaps
 anemia;
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RESULT 6
AAW70384
ID AAW7
XX
AC AAW7
AC AAW7
XX
AC DT 02-I
XX
DE huma
XX
KW Huma
 WAX SX RX RX RX RX RX RX RF F F X RX SOSOSOSOSOSOSOSOSOSOSOSOSOSOSOSOS
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 S
 The present invention provides liquid formulation of antibodies or CC antibody fragments that immunospecifically binds to an interleukin-9 (II-CC 9) polypeptide. The formulation exhibit stability, low to undetectable CC levels of aggregation and very little to no loss of the biological CC activities of antibodies and its fragments, even during the long period of storage. The antibody formulation of the invention is useful for CC preventing, managing, treating or ameliorating a disease or disorder CC prolypeptide, II-9R or one or more subunits, inflammatory disorders such as asthma or an allergy, autoimmune disorders such as diabetes, CC Hashimoto's disease, autoimmune adrenal insufficiency, pure red cell canemia, multiple sclerosis, rheumatoid cardittis, systemic lupus CC syrthematosus, rheumatoid arthritis, chronic inflammation, Sjogren's CC erythematosus, rheumatoid arthritis or scleroderma, proliferative CC disease or an infection (preferably, a respiratory infection). The CC disease or an infection (preferably, a respiratory infection). The CC invention is also useful in gene therapy, chemotherapy and to generate continual disease models. The present sequence is a heavy chain variable CC (VH) domain complementarity determining region, CDR3 peptide present in human interleukin-9 (II-9) antibody 4D4. Note: The present sequence is CC the SEQ ID NO: 3 shown on table 1 in page 2 in disclosure of the sequence listing of the specification (4D4 H2-1 D11, 7F3, 71A10, 7F3 CC 2D3, 7F3com-2H2, 7F3com-3H5 and 7F3com-3D4) (see AED66887).
 Query Match
Best Local S
Matches 5
 Human; regeneration-associated serpin-1 protein; RASP-1; liver disorder;
 New antibody formulation comprises antibody or antibody fragment that immunospecifically binds to an IL-9 polypeptide, useful for preventing, managing, treating, or ameliorating, e.g. asthma, allergy, diabetes, or multiple sclerosis.
 human regeneration-associated serpin-1 peptide
 AAW70384;
 AAW70384 standard; protein; 13
 Sequence 13 AA;
 Disclosure; SEQ ID NO 3; 125pp; English.
 WPI; 2005-796072/81.
 12-APR-2005; 2005US-00105268
 US2005260204-A1
 Homo sapiens
 respiratory-gen.; interleukin-9; cytokine; lymphokine; antibody; heavy chain variable region; 4D4 H2-1 D11; 7F3; 71A10; 7F3 22D3; 7F3com-2H2; 7F3com-3H5; 7F3com-3D4.
 12-APR-2004; 2004US-0561845P
 (MEDI-) MEDIMMUNE INC
 24-NOV-2005
 Similarity 5; Conserv
 YYGSDYVK 10
 YLGEEYVK 11
 Conservative
 (first entry)
 44.1%;
 1;
 Score 30; DB 9;
Pred. No. 6.1e+02;
1; Mismatches 2
 Length 13;
 Indels
 0
 Gaps
 0
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RESULT 7
AEA79013
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 5
 Matches
 Query Match
Best Local (
 This is the amino acid sequence of the human regeneration-associated serpin-1 peptide (RASP-1), used in the method of the invention to develop products for the diagnosis and treatment of liver conditions. The RASP-1 polypeptides are involved in regeneration of the liver and are useful in the diagnosis, prognosis and treatment of liver-associated disorders, such as viral infection, autoimmunity, hepatitis or cirrhosis. They can also be used in methods for maintaining cultured cells or tissues, such as hepatocytes cells or tissues, prior to transplantation. In addition, RASP-1, can be used to promote hepatocyte growth in vitro, in order to, e.g. facilitate production of growth factors, such as hepatocyte growth factors, such as hepatocyte growth factors (HGF), that are produced by them. The regulatory region of RASP-1 allows tissue-specific expression (i.e. liver) of an operatively linked
 New isolated human regeneration-associated serpin-1 - used to develop products for the diagnosis and treatment of liver conditions, e.g. viinfection, autoimmunity, hepatitis or cirrhosis.
 protein quantitation; protease; high-result townstands proteomics; genomics; bioinformatics; Bovine Serum Al
 mass spectrometry; peptide index; protein identification;
 Bovine Serum Albumin indexed peptide database peptide #229.
 11-AUG-2005
 AEA79013 standard; peptide; 13 AA
 Sequence 13 AA;
 Example 1; Page 27; 40pp; English.
 WPI; 1998-446810/38
 Purchio AF;
 03-FEB-1997;
 03-FEB-1998;
 06-AUG-1998.
 Homo sapiens
 hepatocyte growth factor; HGF.
 viral infection; hepatitis; cirrhosis; transplantation;
 WO2003054549-A2
 gs sp
 (HEPA-) HEPATIX INC
 Local Similarity
nes 5; Conserv
 4
 1 YEKYLGEEY 9
 YENQIWEEY 12
 Conservative
 (first entry)
 97US-0036842P
 98WO-US001962
 41.2%;
 ۲.
 Score 28;
Pred. No.
 treatment of liver conditions, e.g. viral
 1.3e+03;
 DB 2;
 Length 13;
 spectrometry;
 <u>.</u>
 to develop
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08-DEC-2001; 2001US-0340460P 09-DEC-2002; 2002WO-GB005571.

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 fragmentation of a polypeptide or protein; determining a first physico-
c chemical property other than mass or mass to charge ratio of the
molecules comprising peptides; and optionally determining a second,
third, fourth and/or fifth physico-chemical property of the molecules
cc comprising peptides; and a mass spectrometer comprising a mass analyser
cc for accurately determining the mass to charge ratio of the first
cc molecules, and means for identifying the first molecules of the basis of
at least the first physico-chemical property and the accurately
cd determined mass to charge ratio of the first molecules and optionally on
the basis of the second, third, fourth and/or fifth physico-chemical
cc property. The method and spectrometer are useful in protein
dentification, protein quantitation, proteases, high-resolution mass
spectrometry, proteomics, genomics and bioinformatics. This sequence
represents a peptide from an indexed peptide database created by the
created and spectrometry method of the invention.
 Query Match
Best Local Similarity
 Matches
 Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
 comprises mass analysing the first molecules in a first mixture and accurately determining the mass to charge ratio of the first molecules in the first mixture. The invention further relates to: generating an index for use in identifying molecules of biological origin by mass spectrometry by accurately determining the masses or mass to charge ratios of molecules comprising peptides resulting from the digestion or
 A method of mass spectrometry, useful in protein identification and quantitation, by mass analyzing the first molecules in the first miand accurately determining the mass to charge ratio of the first
 The invention relates to a novel method of mass spectrometry. The method comprises mass analysing the first molecules in a first mixture and
19-OCT-2000; 2000US-0242350P.
20-APR-2001; 2001US-0285624P.
 HLA binding
 Sequence 13
 Geromanos S,
 18-OCT-2001;
 15-MAY-2003
 10-MAR-2005
 ADW33519 standard; peptide; 13 AA.
 molecules in the first mixture.
 14-MAR-2002; 2002US-0364847P
 WO2003040165-A2
 (MICR-) MICROMASS LTD
 4;
 IGEEYI 6
 LGEEYV 10
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 epitope #4269.
 Conservative
 Fig 9H; 123pp;
 2001WO-US051650
 (first entry)
 Dongre A,
 oy mass analyzing the first molecules in the first determining the mass to charge ratio of the first
 41.2%;
 Opiteck G,
 English.
 Score 28; DB 7;
Pred. No. 1.3e+03;
 Mismatches
 Silva J;
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 Length 13;
 Indels
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 Gaps
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ADW34752

ID ADW3

XX ADW3

XX ADW3

XX ADW3

XX ADW3

XX HLA

XX Viru

XW Viru

XW MHC

XW WO2C

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 Matches
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 The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
 Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
 Sequence 13
 WPI; 2003-441519/41
 10-MAR-2005
 ADW34752 standard; peptide; 13 AA.
 having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
 New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing
 19-OCT-2000; 2000US-0242350P.
20-APR-2001; 2001US-0285624P.
 15-MAY-2003
 WO2003040165-A2
 Unidentified
 HLA binding epitope #5502.
 Claim 1; Page 52-379; 382pp; English.
 binding motifs for
viral diseases and
 18-OCT-2001; 2001WO-US051650
 (EPIM-) EPIMMUNE INC
 N
 EPIMMUNE INC
 4.
 Similarity
 KYVKQNYLK 10
 KYLGEEYVK 11
 Sidney J,
 Sidney J,
 ¥,
 Conservative
 (first entry)
 cancer.
 38.2%;
 Southwood
 composition comprising at least
 ų.
 Score 26; DB 7; Le
Pred. No. 2.7e+03;
 S
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 Length 13;
 Indels
 one peptide
 0,
 Gaps
 0
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RESULT 10
AAW21308
ID AAW21308
AC AAW21
XX AC AAW21
XX COMpe
KW Hydro
KW Compe
KW Charg
KW Hydro
KW Gonad
KW Alzhe
KW Apoli
KW Alzhe
KW Hepat
XX Homo
XX Homo
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 from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cells from the patient with the composition cited above. The patient composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADM29251-ADM37745 represent epitopes of the invention as given in Tables 2-31.
 Matches
 Query Match
Best Local
 Hydrophilic; signal oligopeptide; hydrophilicity maxima; vaccine; SIV; competitive inhibitor; feedback regulator; synthesis; gastrin precursor; charge; polarity; farnesyl synthetase; plasminogen activator inhibitor 1 hydroxymethylglutaryl coenzyme A reductase; glucagon precursor; rhesus; gonadoliberin precursor; plasminogen activator inhibitor 2; prorenin; Alzheimer amyloid A4; corticotropin releasing factor binding protein; apolipoprotein E; herpes virus 1 glycoprotein B; HSV1; human; OMVVS; herpes virus 2 glycoprotein B; HSV2; collagenase; apolipoprotein A; Treponema pallidum membrane protein; TMPA; islet amyloid polypeptide; fibroblast MMPI; schistosoma elastase precursor; schistosomin; hepatitis delta antigen; rev protein; HIV; VILV; angiotensinogen.
 Glucagon precursor derived signal oligopeptide #13.
 Claim 1; Page 52-379; 382pp; English.
 New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
 WPI; 1995-263953/34
 Rath M;
 29-JUL-1997 (first entry)
 AAW21308;
 Sequence 13 AA;
 14-JAN-1994;
 12-JAN-1995;
 WO9519568-A1
 AAW21308 standard; peptide; 13
 The invention relates to a composition comprising at least one peptide
 Homo sapiens
 N
 3 KYLGEEYVK 11
 4.
 Similarity
 KYVKQNYLK 10
 Conservative
 94US-00182248
 95WO-US000575.
 38.2%;
 Score 26; DB
Pred. No. 2.7e
3; Mismatches
 DB 7; 1
2.7e+03;
 2; Indels
 Length 13;
 0,
 Gaps
 0
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Identifying signal oligopeptide(s) in protein sequence(s) - shown as regions of max. hydrophilicity, used in modulating communication between protein(s).
 Claim 5; Page 40; 88pp; English
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The sequences given in AAW21201-560 represent hydrophilic signal oligocc paptides. These signal oligopeptides are localised on the surface of the
cc protein and are represented by the hydrophilicity maxima of the protein.
cc protein and are represented by the hydrophilicity maxima of the protein.
cc protein and are represented by the hydrophilicity maxima of the protein.
cc protein and are represented by the hydrophilicity maxima of the protein.
cc protein are represented by the hydrophilicity maxima of the protein and charge consideration of conformation and charge contains the signal sequence. These oligopeptides may be used as vaccines continuously the metabolic action or interaction of a selected protein by contain gits specific signal sequences, as competitive inhibitors to prevent conception as feedback regulators to reduce synthesis rate of a selected concition as feedback regulators to reduce synthesis rate of a selected concition. These peptides may be modified by omitting one or more amino calds at the N- and/or C-terminal, by substituting one or more amino calds without consideration of charge and polarity, by substituting one or more amino acids with amino acid residues with similar charge and/or the polarity by onitring one or more amino acids with similar charge and/or the protein. by omitting one or more amino acids or a combination of these

Query Match Best Local ( Matches l Similarity 5; Conserve N YSKYLDNRRAK 12 YEKYLGEEYVK 11 Conservative 36.8%; 0 Score 25; Pred. No. Mismatches DB .9e+03; 6, Length 13, Indels 0

0

Sequence 13 AA;

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29-JUL-1997 AAW21301 standard; peptide; 13 (first entry)

Glucagon precursor derived signal oligopeptide #6.

RESULT 11
AAW21301
ID AAW21301
AC AAW21
AC AAW21
AC AAW21
AC AAW21
AC Gluca
AC Gluca
AC Charg
CKW Compe
KW Charg
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A Hydrophilic; signal oligopeptide; hydrophilicity maxima; vaccine; SIV; competitive inhibitor; feedback regulator; synthesis; gastrin precursor; charge; polarity; farnesyl synthetase; plasminogen activator inhibitor 1, hydroxymethylglutaryl coenzyme A reductase; glucagon precursor; rhesus; gonadoliberin precursor; plasminogen activator inhibitor 2; prorenin; Alzheimer amyloid A4; corticotropin releasing factor binding protein; apolipoprotein E; herpes virus 1 glycoprotein B; HSV1; human; OMVVS; herpes virus 2 glycoprotein B; HSV2; collaganase; apolipoprotein A; Treponema pallidum membrane protein; TMPA; islet amyloid polypeptide; fibroblast MMP1; achistosoma elastase precursor; schistosomin; hepatitis delta antigen; rev protein; HIV; VILV; angiotensinogen.

12-JAN-1995; WO9519568-A1 Homo sapiens 95WO-US000575

(RATH/) RATH

14-JAN-1994;

94US-00182248

WPI; 1995-263953/34.

Identifying signal oligopeptide(s) in protein sequence(s) - shown as

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ARESULT 12
AAW21133
ID AAW21
AC AAW21
AC AAW21
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 The sequences given in AAW21201-560 represent hydrophilic signal oligopeptides. These signal oligopeptides are localised on the surface of the protein and are represented by the hydrophilicity maxima of the protein. These peptides are enriched in charged amino acids arranged with neutral spacer amino acids. The specific signal character of these oligopeptides is determined by a characteristic combination of conformation and charge within the signal sequence. These oligopeptides may be used as vaccines in the treatment of human disease, as competitive inhibitors to prevent or reduce the metabolic action or interaction of a selected protein by blocking its specific signal sequences, or as therapeutic agents to blocking its specific signal sequences, or as therapeutic agents to function as feedback regulators to reduce synthesis rate of a selected protein. These peptides may be modified by omitting one or more amino acids at the N- and/or C-terminal, by substituting one or more amino acids with amino acid residues with similar charge and/or polarity, by omitting one or more amino acids with amino acid residues with similar charge and/or polarity, by omitting one or more amino acids or a combination of these
 Matches
 Best Local Similarity
 Query Match
 hydroxymethylglutaryl coenzyme A reductase; glucagon precursor; rhesus; gonadoliberin precursor; plasminogen activator inhibitor 2; prorenin; Alzheimer amyloid A4; corticotropin releasing factor binding protein; apolipoprotein E; herpes virus 1 glycoprotein B; HSV1; human; OMVV5; herpes virus 2 glycoprotein B; HSV2; collagenase; apolipoprotein A; Treponema pallidum membrane protein; TMPA; islet amyloid polypeptide; fibroblast MMP1; schistosoma elastase precursor; schistosomin; hepatitis delta antigen; rev protein; HIV; VIIV; angiotensinogen.
 Hydrophilic; signal oligopeptide; hydrophilicity maxima; vaccine; SIV; competitive inhibitor; feedback regulator; synthesis; gastrin precursor; charge; polarity; farnesyl synthetase; plasminogen activator inhibitor 1
 Glucagon precursor derived signal oligopeptide #18.
 AAW21313 standard; peptide; 13 AA
Identifying signal oligopeptide(s) in protein sequence(s) - shown as regions of max. hydrophilicity, used in modulating communication bet
 14-JAN-1994;
 12-JAN-1995;
 20-JUL-1995.
 29-JUL-1997
 AAW21313;
 Sequence 13
 Claim 5; Page 39; 88pp; English.
 protein(s)
 WPI; 1995-263953/34.
 WO9519568-A1
 Homo sapiens
 regions of max. hydrophilicity, used in modulating communication between
 (RATH/) RATH
 N
 س
 <u>ა</u>
 YEKYLGEEYVK 11
 YSKYLDNRRAK 12
 Conservative
 Š
 (first entry)
 94US-00182248
 95WO-US000575.
 36.8%;
45.5%;
 0; Mismatches
 Score 25; DB 2;
Pred. No. 3.9e+03;
 6;
 Length 13;
 Indels
 0;
 Gaps
 0
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communication between

WPI; 2004-517671/49

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 The sequences given in AAW21201-560 represent hydrophilic signal oligocopeptides. These signal oligopeptides are localised on the surface of the protein and are represented by the hydrophilicity maxima of the protein. These peptides are enriched in charged amino acids arranged with neutral spacer amino acids. The specific signal character of these oligopeptides government of a characteristic combination of conformation and charge within the signal sequence. These oligopeptides may be used as vaccines in the treatment of human disease, as competitive inhibitors to prevent correduce the metabolic action or interaction of a selected protein by blocking its specific signal sequences, or as therapeutic agents to conform as feedback regulators to reduce synthesis rate of a selected protein. These peptides may be modified by omitting one or more amino acids at the N- and/or C-terminal, by substituting one or more amino acids without consideration of charge and polarity, by substituting and/or correct amino acids with amino acid residues with similar charge and/or conformation and charge and polarity.
 Matches
 Query Match
Best Local
 peripheral nervous system disorder; postoperative nerve damage; traumatic nerve damage; hipaired myelination; post-ischaemic damage; stroke; Parkinson's disease; Alzheimer's disease; Huntington's disease; dementia; multiinfarct dementia; sclerosis; nerve degeneration; diabetes mellitus; schizophrenia; mood disorder; manic depression; neuro-muscular connection; organ transplantation; genetic atrophic muscle disorder; traumatic atrophic muscle disorder; degenerative condition; acute myocardial infarction; angiogenesis;
 receptor binding site; neuroprotective; vulnerary; antidiabetic; nephrotropic; gene therapy; fibroblast growth factor receptor; FGFR receptor ligand; neural cell adhesion molecule; NCAM; solid tumour; neoanglogenesis; central nervous system disorder;
 Berezin V,
 Synthetic
 20-DEC-2002; 2002DK-00001982
03-MAR-2003; 2003DK-00000330
 18-DEC-2003; 2003WO-DK000901.
 protein interaction modulation; cell-surface receptor;
 Putative fat-like cadherin precursor FGFR binding motif peptide SEQ:32
 23-SEP-2004 (first entry)
 ADP83124;
 ADP83124 standard; peptide; 13
 Sequence 13
 WO2004056865-A2
 wound-healing; revascularisation
 polarity, by omitting one or more amino acids or a combination of these
 Claim 5; Page 41; 88pp; English.
 protein(s).
 (ENKA-) ENKAM PHARM AS
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 1 YEKYLGEEYVK 11
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 Similarity
 YSKYLDNRRTK 12
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 Conservative
 36.8%;
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 Bock E;
 Score 25; DB 2; Le
Pred. No. 3.9e+03;
 Length 13;
 Indels
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 FGFR;
 Gaps
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 cc between at least two different proteins, where one of the proteins is care variant or by a polypeptide having a binding site to the receptor, where cat least a part of the binding site comprises at least one of the greent or cat least a part of the binding site comprises at least one of the sequences given in SEQ ID NO:1 to 146 (ADP83093 to ADP83238), or comprise the form of the sequences given in SEQ ID NO:1 to 146 (ADP83093 to ADP83238), or companies of the homologues, comprising (i) providing a compound cat step (i) to at least two different proteins. The compound has neuroprotective, vulnerary, antidiabetic and nephrotropic activities, and can be used in gene therapy. The method is useful in modulating the control is represented by a functional cell-surface receptor, selected can be used in gene therapy. The method is useful in modulating the compound of step (i) to at least two different proteins. The compound has neuroprotective, vulnerary, antidiabetic and nephrotropic activities, and can be used in gene therapy. The method is useful in modulating the interaction between at least two different proteins, where one of the control is represented by a functional cell-surface receptor, selected from the family of fibroblast growth factor receptors (FGFRs), and the control is represented by a functional cell-surface receptor, selected cather protein is the receptor ligand, can be used for treating normal, cells; solid tumour requiring neoanglogenesis; diseases and conditions of the central and peripheral nervous system, of the muscles or of various of the central and peripheral nervous system, of the muscles or of various of the central method. Parkinson's disease, Alzheimer's disease, e.g. currently dementia, sclerosis, nerve degeneration associated with diabetes mellitus, disorders affecting the circadian clock or neuro-muscular transmission, and schizopherala.
 Query Match
 nerve degeneration associated with glabetes merricus, unsuccession, the circadian clock or neuro-muscular transmission, and schizophrenia, mood disorders, such as manic depression; diseases or conditions of the muscles including conditions with impaired function of neuro-muscular connections, such as after organ transplantation, or such as genetic or transmit atrophic muscle disorders; degenerative conditions of the gonada, of the pancreas such as diabetes mellitus type I and II, of the kidney such as nephrosis or of the heart, liver and bowel; for preventing cell death of heart muscle cells, such as after acute myocardial infarction, or after angiogenesis; for promoting wound-healing; for revascularisation; or for stimulating the ability to learn and/or the
Respiratory infection; human; interleukin-9 antagonist; IL-9; respiratory syncytial virus; anti-RSV antigen antibody; leukotriene modifier; asthma-like symptom; wheezing; allergy;
 Modulating the interaction between at least two different proteins by providing a compound capable of interfering with the receptor and the polypeptide interaction and presenting the compound to the different
 Human 4D4com-XF-9 variable heavy chain (VH)
 ADU07237
 ADU07237 standard;
 Sequence 13
 Claim 45; SEQ ID NO 32; 154pp; English.
 Local
 Similarity
5; Conserv
 EKYLGEEYVK 11
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 Conservative
 (first entry)
 peptide;
 36.8%;
 describes a method for modulating the interaction
 13
 Score 25; DB 8;
Pred. No. 3.9e+03;
1; Mismatches 4
 CDR3.
 Length 13;
 Indels
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 Gaps
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immunomodulatory; anti-inflammatory; anti-viral; antibiotic; antifungal; mast cell modulator; virucide; antibacterial; fungicide; antiasthmatic; antiallergic; 4D4com-XF-9 antibody; variable heavy chain; VH; complementarity determining region; CDR.
 WPI; 2004-775524/76.
 11-APR-2003; 2003US-0462307P.
10-JUN-2003; 2003US-0477801P.
 WO2004091519-A2
 (MEDI-) MEDIMMUNE INC
 12-APR-2004; 2004WO-US011329
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Managing, treating or ameliorating in a human subject a respiratory infection or its symptoms caused by bacteria, fungi or respiratory syncytial virus by administering an IL-9 antagonist.

Disclosure; SEQ ID NO 12; 276pp; English

Crespiratory infection or its symptom in a human subject by administering a crespiratory infection or its symptom in a human subject by administering can interleukin-9 (IL-9) antagonist. The method further comprises can anti-respiratory syncytial virus (anti-key) antigen antibody. The anti-RSV antigen antibody is palivizumab. The method further comprises administering a leukorriene modifier, which is method of preventing the development, onset or progression of asthma or one or more asthma-like symptoms in a child, a method of preventing, cansaing, treating or ameliorating wheezing in a pre-term infant, infant or child, a method of preventing, managing, treating or ameliorating method of preventing, managing, treating or method of preventing, managing, treating or method of preventing, managing, treating or method rection comprises administering an IL-9 antagonist.

Crespiratory infection comprises administering an IL-9 antagonist. The comprises administering an IL-9 antagonist. The contaminatory agent, an anti-viral agent, an anti-breapy that is not administering wheezing in a human subject comprises administering an IL-9 antagonist. The crifical points and at least one other therapy that is not administering to the subject a nati-transming-doublin E (IgE) antibody, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, an anti-brica, and anti-brica, an anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-brica, and anti-br

Sequence 13 AA;

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 Matches
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 Similarity
4; Conser
 Conservative
 36.8%;
 Score 25; I
Pred. No. 3
 Mismatches
 DB 8;
 Length 13
 <u>.</u>.
 Gaps
 0
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Db 3 YYGSDHVK 10
RESULT 15
ADT88304

ADT88304 standard; peptide; 13 AA.

ADT88304;

13-JAN-2005 (first entry)

Human IL-19 antibody variable heavy domain CDR3 seqid 12.

respiratory; antiinflammatory; immunosuppressive; antiasthmatic; antiallergic; antiarthritic; neuroprotective; antirheumatic; immunotoxin; angiogenesis inhibitor; TNF alpha antagonist; IL-9 antagonist; IL-9; IL-9 antagonist; AD4; AD4 H2-1 D11; AD4com-XF9; AD4com-2F9; 7F3; 71A10; 22D3; 7F3com-2H2; 7F3com-3H5; 7F3com-3D4; IL-9 associated disorder; respiratory infection; respiratory syncytial virus; parainfluenza virus; human metapneumovirus; inflammatory disorder; asthma; allergy; arthritis; autoimmune disorder; rheumatori disorder; asthmatis; multiple sclerosis; variable heavy domain; complementarity determining region 3; CDR3.

Homo sapiens

WO2004091510-A2.

28-OCT-2004.

12-APR-2004; 2004WO-US011172.

11-APR-2003; 2003US-0462259P. 10-JUN-2003; 2003US-0477797P.

(MEDI-) MEDIMMUNE INC.

Reed JL;

WPI; 2004-775520/76.

New IL-9 antibody comprising a variable heavy or light domain or a complementarity determining region having an amino acid sequence of e.g., 4D4, useful in treating or preventing e.g., respiratory infection or inflammatory disorder.

Disclosure; SEQ ID NO 12; 291pp; English.

CC (WH) or variable light (VL) domain having an amino acid sequence of the CV (WH) or variable light (VL) domain having an amino acid sequence of the CC (WH) or VL domain of 404, 404 H2-1 DI1, 4D4com-XF9, 4D4com-ZF9, 7F3, 71A10, CC 22D3, 7F3ccm-2H2, 7F3com-3H5 or 7F3com-3D4; or a complementarity CC determining region (CDR) having an amino acid sequence of a CDR of 4D4, 4D4 H2-1 DI1, 4D4com-XF9, 4D4com-ZF9, 7F3, 71A10, 22D3, 7F3com-2H2, CC 7F3com-3H5 or 7F3com-3D4. The antibody immunospecifically binds to a CC comprising the II-9 antibody and a carrier; a method for preventing, cc treating, managing or ameliorating a disease or disorder characterised by aberrant expression or activity of an II-9 polypeptide or an II-9R; a cc method of diagnosing, prognosing, or monitoring a disorder or disease cc characterised by aberrant expression or activity of an II-9R; a kit comprising the CC or aberrant expression or activity of an II-9R; a kit comprising the CC antibody and instructions for use, in one or more containers; and an CC article of manufacture comprising a packaging material, where the pharmaceutical agent comprises the II-9 antibody and a carrier. The II-9 antibody is useful in preparing a composition for diagnosing, treating or preventing a disease or disorder characterised by aberrant expression or activity of an II-9 polypeptide or an II-9 polypeptide or antibody is useful in preparing a composition for diagnosing, treating or preventing a disease or disorder characterised by aberrant expression or cc activity of an II-9 polypeptide or an II-9R, e.g.; respiratory infection CC caused by bacteria or respiratory syncytial virus, parainfluenza virus or human metapneumovirus, inflammatory disorder such as asthma, allergy or arthritis or autoimmune disorder such as rheumatoid arthritis and continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the continue of the con

antibody variable heavy domain complementarity determining region

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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-13 <LE5>
 A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-13 <LE3>
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A; Residues: 1-13 < LE2>
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C;Date: 06-Jan-1995 #sequence
C;Accession: $47361
R;Lehner, P.J.
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 A; Cross-references: UNIPARC: UPI0000116673;
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 A; Molecule type: mRNA
A; Residues: 1-13 < LEH>
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 A; Accession: S47379
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 C; Keywords: T-cell receptor
 A; Description: Human HLA-A0201
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 ;Keywords:
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 Molecule type: mRNA
Residues: 1-13 <LE6>
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 Cross-references: UNIPARC:UPI000011667B; EMBL:Z35685; NID:g527459; PIDN:CAA84754.1;
 Status: preliminary
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 EMBL: 235708; NID: 9527509;
 EMBL: 235675; NID: 9527529;
 EMBL: 235674; NID: g527527;
 EMBL: Z35700; NID: g527493;
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C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: $47383
R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognit
**Poference number: $47355
 RESULT 6
$47372
T-cell antigen receptor VJ junction beta chain -
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-199
C;Accession: $47372
C;Accession: $47372
R;Lehner, P.J.
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C:Date: 06-Jan-1995 #sequence_revision 06-Jan-199
C:Accession: $47368
A;Molecule type: mRNA
A;Restdues: 1-13 <LEHS
A;Cross-references: UNIPARC:UPI000011668D; EMBL:Z35709; NID:g527513; PIDN:CAA84778.1; PII
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 A;Modecule type: mRNA
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A;Residues: 1-13 <LEH>
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A;Reference number: S47355
 submitted to the EMBL Data Library, August 1994 A;Description: Human HLA-A0201 restricted recognition A;Reference number: S47355 A;Accession: S47368
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 submitted to the EMBL Data Library, August 1994 A;Description: Human HLA-A0201 restricted recogn A;Reference number: $47355 A;Accession: $47384
 C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: S47384
R;Lehner, P.J.
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A;Residues: 1-13 <LEH>
 T-cell antigen receptor VJ junction beta chain - human (fragment)
(;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence revision 25-Oct-1996 #text_change 05-Nov-1999
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 A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-13 <LEH>
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 A;Cross-references: UNIPARC C;Keywords: T-cell receptor
 A; Molecule type: mRNA
A; Residues: 1-13 <LE2>
 submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition of influenza
 R;Lehner, P.J. submitted to the EMBL Data Library,
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 T-cell antigen receptor VJ junction beta chain -
 S47384
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 A;Cross-references: UNIPARC:UPI0000116674; EMBL:Z35699; NID:g527491; PIDN:CAA84768.1;
 A; Accession: S47374
 A; Reference number: S47355
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 #text_change 09-Jun-2000
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J. Protein Chem. 10, 415-425, 1991
A;Title: Clostridium botulinum types A, B, C1, and E produce proteins with or without he A;Reference number: A44644; MUID:92143938; PMID:1781887
A;Contents: type B
A;Accession: G44644
 neurotoxin-associated protein type B Hn+ 35K chain, C;Species: Clostridium botulinum C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994
 C;Accession: A32734

R;Micanovic, R:; Ray, P:; Kruggel, W:; Lewis, R.V.
Biochem. Biophys. Res. Commun. 118, 299-303, 1984

A;Title: Purification and sequence of an opioid peptide
A;Reference number: A32734; MUID:84128045; PMID:6546517
 A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-13 <SOM>
A;Cross-references: UNIPROT:Q9R5N5; UNIPARC:UPI00000B9B79
A;Note: sequence extracted from NCBI backbone (NCBIP:83785)
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A;Residues: 1-13 <LIM>
 Southeast Asian J. Trop. Med. Public Health 21, 388-396, 1990 A;Title: Characterization of a Plasmodium falciparum epitope : A;Reference number: A60396; MUID:91164876; PMID:1706114 A;Accession: E60396
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 A;Molecule type: protein
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A;Cross-references: UNIPROT:Q7M2Z6; UNIPARC:UPI00001765B8
C;Superfamily: proenkephalin
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 R;Somers, E.; DasGupta, B.R.
 C; Accession: G44644
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 C; Keywords: neuropeptide;
 A; Accession: A32734
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 A;Cross-references: UNIPARC:UPI000017B5EF; GB:M31305
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 ;Keywords: hemagglutinin
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Pred. No. 7.7e+03;
3; Mismatches 4
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Pred. No. 5.2e+03
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T-cell antigen receptor VJ junction beta chain - human
C.Species: Homo sapiens (man)
C.Species: Homo sapiens (man)
C.Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 05-Nov-1999
C.Accession: S47373
R.Lehner, P.J.
Submitted to the EMBL Data Library, August 1994
A.Description: Human HLA-A0201 restricted recognition of influenza A is dominated by T
A.Reference number: S47373
A.Reference number: S47355
A.Accession: Settinary
A.Accession: Settinary
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(;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
(;Accession: PT0290
R,Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0290
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J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-13 <YAM'>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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 Ig heavy chain CRD3 region (clone 2-121B) - human (fragment)
c;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0263
 A;Cross-references: UNIPARC:UPI0000116671; EMBL:Z35672; NID:g527489; PIDN:CAA84741.1; PIC:Keywords: T-cell receptor
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13 | Q8WEJB GNEGN<br>Q83171 9VIRU<br>Q5EX74 SPHPU<br>Q5EX79 9HIVI<br>NEUT CAVPO<br>Q2VRYO 9PEZI<br>Q2VRY1 9PEZI<br>Q2VRY2 9PEZI<br>Q2VRY3 9PEZI<br>Q2VRY3 9PEZI<br>Q2VRY4 9PEZI<br>Q2VRY4 9PEZI<br>Q2VRY6 9PEZI<br>Q2VRY6 9PEZI<br>Q2VRY6 9PEZI<br>Q2VRY6 9PEZI<br>Q3S793 YEAST<br>Q49NC3 9PEZI<br>Q49NC3 9PEZI                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
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| 3 AA.<br>Swiss-Prot.<br>Vertebrata;<br>a; Ranoidea;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | DB 2; Length 13<br>1.3e+04;<br>.ches 1; Indels        | latabas<br>0://www<br>ion-No<br>.RC64;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | 01-NOV-1996, integrated into UniProtKB/TrEMBL. 01-NOV-1996, sequence version 1. 07-FEB-2006, entry version 14. 072661. Wycobacterium leprae. Bacteria; Actinobacteridae; Actinomycet Corynebacterinee; Mycobacteriaceae; Mycobacterium. NCBI_TaxID=1769; [1] NUCLEOTIDE SEQUENCE. Smith D.R.; Smith D.R.; Smith D.R.; Submitted (APR-1995) to the EMBL/GenBank/DDBJ databases                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | NTS                  | 20                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |
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RESULT 3
Q654R5_LEICH
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RESULT 4

Q3LYXO 9INFA
ID Q3LYXO.9INFA PRELIMINARY; PRT; 13 AA.
AC Q3LYXO;
DT 25-OCT-2005, integrated into UniProtKB/TrEMBL
DT 25-OCT-2005, sequence version 1.
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 "Temporins, a
temporaria.";
Eur. J. Bioch
 STRAIN=MHOM/BR/00/1669;
PubMed=15888291; DOI=10.1016/j.exppara.2005.02.008;
Purdy J.E., Donelaon J.E., Wilson M.E.;
"Leishmania chagasi: The alpha-tubulin intercoding region results in constant levels of mRNA abundance despite protein synthesis inhibition
 MOD_RES
 -I- FUNCTION: Has antibacterial activity against Gram-
Gram-positive bacteria.
-I- SUBCELLULAR LOCATION: Secreted protein.
-I- TISSUE SPECIFICITY: Expressed by the skin glands.
-I- SIMILARITY: Belongs to the frog skin active pepti-
Brevinin subfamily.
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 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae; Leishmania.
NCBI TaxID=44271;
 05-JUL-2004, integrated into UniProtKB/TrEMBL 05-JUL-2004, sequence version 1.
 EMBL; AY466450; AAR88132.1; -; Genomic_DNA.
 NUCLEOTIDE SEQUENCE.
 Alpha tubulin (Fragment).
 07-FEB-2006, entry version
 Amidation; Amphibian defense peptide; Antibiotic; Antimicrobial;
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 growth state."
 Parasitol. 110:102-107(2005).
 J. Biochem. 242:788-792(1996).
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2; Mismatches 1
 3F361B06B9741B18 CRC64;
 Score 21; DB 1; Length 13; Pred. No. 2.9e+04;
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/FTId=PRO_0000043585.
Leucine amide.
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DE UDP-galactose-
GN Name=GAL10;
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 Kanai T., Ueda M., Tanaka A.; "Candida tropicalis promoter region of GAL1 and GAL10 gene."; Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
 Eukaryota; Fungi; Ascomycota; Saccharomycetales; mitosporic NCBI TaxID=5482;
 01-MAY-1999, integrated into UniProtKB/TrEMBL 01-MAY-1999, sequence version 1. 07-FBB-2006, entry version 13. UDP-galactose-4-epimerase (Fragment).
 Viral ma
 EMBL;
 STRAIN-A/Shorebird/Delaware/275/2001;
PubMed=16081121; DOI=10.1016/j.virol.2005.07.003;
Ilyushina N.A., Govorkova E.A., Webster R.G.;
"Detection of amantadine-resistant variants among viruses isolated in North America and Asia.";
Virology 341:102-106(2005).
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Candida tropicalis (Yeast).
 SEQUENCE
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Name=M;
Influenza A virus (A/shorebird/Delaware/275/2001(H9N7)).
Viruses; ssRNA negative-strand viruses; Orthomyxoviridae;
Influenzavirus A.
 07-FEB-2006, entry version 5. Matrix protein 1 (Fragment).
 NUCLEOTIDE SEQUENCE.
 NUCLEOTIDE SEQUENCE.
 NCBI_TaxID=342210;
 matrix protein.
 DQ107506; ABA12424.1; -; mRNA.
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 Score 20; DB 2;
Pred. No. 4.2e+04;
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 0B559D6AE18BC720 CRC64;
 274716F5CDFBC2C8 CRC64;
 Saccharomycotina; Saccharomycetes; Saccharomycetales; Candida.
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Q86129_9RHAB
Q86129;
 "Structure of the gene N: gene NS intercistronic junction in the genome of vesicular stomatitis virus."; Cell 17:673-681(1979).
 01-NOV-1996, sequence version 07-FEB-2006, entry version 16. NS protein (Fragment). Vesicular stomatitis virus.
 Acyl-protein synthetase Name=luxE;
 Q3KYR7;
08-NOV-2005, integrated into UniProtKB/TrEMBL.
08-NOV-2005, sequence version 1.
07-FEB-2006, entry version 3.
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 Viruses; ssRNA negative-strand viruses; Mononegavirales; Rhabdoviridae; Dimarhabdovirus supergroup; Vesiculovirus
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Photobacterium phosphoreum species group."; Environ. Microbiol. 7:1641-1654(2005).
 Bacteria; Proteobacteria; Gam
Vibrionaceae; Photobacterium.
 SEQUENCE
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 EMBL;
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 EMBL;
 Ast J.C., Dunlap P.V.; "Phylogenetic resolution and habitat specificity of members
 STRAIN=FS-4.2
 Photobacterium phosphoreum.
 NUCLEOTIDE SEQUENCE.
 NCBI_TaxID=11276;
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 PubMed=16156737; DOI=10.1111/j.1462-2920.2005.00859.x;
 NUCLEOTIDE SEQUENCE.
 MEDLINE=80001959; PubMed=89911; DOI=10.1016/0092-8674(79)90274-5;
 V01210; CAA24522.1; -; Genomic_RNA.
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 AY849494; AAX51447.1; -; Genomic_DNA.
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 1556 MW;
 27.9%;
75.0%;
 27.9%;
 (Fragment)
 Gammaproteobacteria; Vibrionales;
 Score 19; DB
Pred. No. 6.16
2; Mismatches
 Score 19; DB
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SEQUENCE 13 AA; 13
 15-DEC-2003, integrated into Unil
15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 8.
Enkephalin precursor (Fragment).
Ovis aries (Sheep).
 EMBL; CAAI01003487; CAI00692.1; -; Genomic_DNA.
Hypothetical protein.
SEQUENCE 13 AA; 1620 MW; F4F2838E31621051 C
 Berriman M., Florens L., Janssen C.S., Pain A., Christophides James K., Rutherford K., Harris B., Harris D., Churcher C.M., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J., Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III, Kafatos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Waters
 05-JUL-2005, integrated int
05-JUL-2005, sequence versi
07-FBB-2006, entry version
Hypothetical protein.
ORFNames=PB400005.00.0;
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 MEDLINE=84128045; PubMed=6546517; Micanovic R., Ray P., Kruggel W., "Purification and sequence of an oproenkephalin.";
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae; Caprinae; Ovis.
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 "A comprehensive survey of the Plasmodium life cycle transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
-i- CAUTION: The sequence shown here is derived from EMBL/GenBank/DDBJ whole genome shotgun (WGS) enti
 NUCLEOTIDE SEQUENCE.

PubMed=15637271; DOI=10.1126/science.1103717;
Hall N., Karras M., Raine J.D., Carlton J.M.,
Parent M. Placone I. Janesen C. S. Dain A.
 Q4YMR8_PLABE
Q4YMR8;
 NCBI_TaxID=9940;
 Sinden R.S.;
 NCBI_TaxID=5821;
 Plasmodium berghei.
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium
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 Biochem.
 preliminary data.
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 Biophys. Res. Commun. 118:299-303 (1984)
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RESULT 11

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OX NCBI_TaxID=573;
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RC STRAIN=M5a1;
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RA Albrecht C., Kleiner D.;
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DR EMBL; Y17587; CAA76799.1; -; Genomic_DNA.
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07-FEB-2006, entry version 8.
Interleukin 1 alpha (Fragment).
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 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Klebsiella.
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 NUCLEOTIDE SEQUENCE.
MEDLINE=99397042; PubMed=10467711;
Maddox J.F., Hawken R.J., Matthew P., Davies K.P.;
"Single strand conformational polymorphisms (SSCPs) in and IL6 genes.";
 Ovis aries (Sheep).
 NCBI_TaxID=9940;
 01-MAY-2000,
 Genet. 30:317-318(1999).
 AF117652; AAD25050.1; -;
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 PubMed=15956661; DOI=10.1534/genetics.105.045120; Borge T., Webster M.T., Andersson G., Saetre G.-P.; "Contrasting Patterns of Polymorphism and Divergence Chromosome and Autosomes in Two Ficedula Flycatcher S Genetics 171:1861-1873(2005).
 OFRSN5_CLOBO PRELIMINARY; PRT; 13 AA. 09RSN5_
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01-MAY-2000, integrated into UniProtKB/TrEMBL
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Clostridium botulinum.
 Picedula albicollis (Collared flycatcher)
Eukaryota; Metazoa; Chordata; Craniata; V
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07-FEB-2006, entry version 3.
Chromo-helicase-DNA-binding on the Z chromoson
 EMBL; AJ869811;
GO; GO:0003677;
 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
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 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 without hemagglutinating activity:
sequences and genes?";
J. Protein Chem. 10:415-425(1991).
 PROTEIN SEQUENCE.
MEDLINE=92143938; PubMed=1781887;
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 Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
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 Craniata; Vertel; Passeriformes;
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IEA.
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 13 AA.
 8.9e+04;
 Vertebrata; Euteleostomi;
ormes; Muscicapidae; Ficed
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 .9e+04;
 Length 13;
 Length 13;
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SOFFR

GO; GO:0004386; F:helicase activity; IEA.

NON TER NON TER SEQUENCE

13

13 AA; 1703 MW;

5FEBFF288DB81B53 CRC64;

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RESULT 14

RESULT 14

RE121 LITRU

ID EI21 LI

AC P82097

DT 23-JAN

DT 01-MAY

DT 01-MAY

DT 01-MAY

OC Electri

OC Pelodr

OC Pelodr

OC Pelodr

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CC Distri

CC Distri

CC Amidat

FT PEPTIC

FT MOD_RE

SQ SEQUEN
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E820-8;
P820-98;
23-JAN-2002, integrated into UniProtKB/Swiss-Prot.
01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 22.
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 EI21 LI7
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
 Aust. J. Chem. 52:639-645(1999).
-!- SUBCELLULAR LOCATION: Secreted protein.
-!- TISSUE SPECIFICITY: Expressed by the skin glands.
 Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.; "Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
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 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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 EYLK 6
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Proline amide.
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TISSUE=Skin secretion;

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 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
 Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.; "Peptides from the skin glands of the Australian buzzing Litori electrica. Comparison with the skin peptides from
 Amidation; Amphibian defense peptide; Direct protein sequencing. PEPTIDE 1 13 Electrin-2.2.
 -!- SUBCELLULAR LOCATION: Secreted protein.-!- TISSUE SPECIFICITY: Expressed by the skin glands.
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 Total number of hits satisfying chosen parameters:
 Searched:
 Scoring table:
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 Perfect score:
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 Score
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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 Gapop 10.0 , Gapext 0
 YEKYLGEEYVKAV 13
 GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd
 Length
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 1 0%
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: 45 summaries
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US-09-15-737-135
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US-09-060-480-7
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 Description
 RESULT 2
US-08-465-325-135
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 RESULT 1
 US-09-534-717-494
 Matches
 GENERAL INFORMATION:
 APPLICANT: Magainin Pharmaceuticals Inc.
APPLICANT: 5110 Campus Drive
APPLICANT: 5110 Campus Drive
APPLICANT: Plymouth Meeting, PA 19462
TITLE OF INVENTION: Biologically Active Peptides Having
TITLE OF INVENTION: N-Terminal Substitutions
NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS:
 COMPUTER READABLE FORM:
 STATE: L
COUNTRY:
ZIP: 200
 STREET: 1300 I. (CITY: Washington
 ADDRESSEE: Finnegan,
ADDRESSEE: Dunner
 20005-3315
 4.
 D.C
 E: Dunner
1300 I. Street,
 USA
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| 29.4             | 29.4             | 29.4            | 29.4              | 29.4              | 29.4            | 30.9               | 30.9              | 30.9              | 30.9             | 30.9             | 30.9            | 30.9              | 30.9              | 30.9             | 30.9             | 30.9             | 30.9              |
| 13               | 13               | 13              | 13                | 13                | 13              | 13                 | 13                | 13                | 13               | 13               | 13              | 13                | 13                | 13               | 13               | 13               | 13                |
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| US-09-126-420A-8 | US-08-844-031-36 | US-08-430-521-9 | US-08-766-725A-17 | US-07-932-455A-17 | US-07-632-027-5 | PCT-US94-10257A-26 | US-09-820-053A-78 | US-09-936-885A-26 | US-09-756-247-35 | US-09-756-247-14 | US-09-379-297-4 | US-09-539-879A-48 | US-09-539-879A-47 | US-09-319-730-10 | US-08-895-590-90 | US-08-895-590-89 | US-08-374-077C-48 |
| Sequence         | Sequence         | Sequence        | Sequence          | Sequence          | Sequence        | Sequence           | Seguence          | Sequence          | Sequence         | Sequence         | Sequence        | Sequence          | Sequence          | Sequence         | Sequence         | Sequence         | Sequence          |
| 8                |                  | 9               | 17,               | 17,               | 'n              | 26,                | 78,               | 26,               | 35,              | 14,              | 4,              | 48,               | 47,               | 10,              | 90,              | 89,              | 48,               |
| Appli            | Appl             | Appli           | Appl              | Appl              | Appli           | Appl               | Appl              | Appl              | Appl             | Appl             | Appli           | Appl              | Appl              | Appl             | Appl             | App              | App               |

ALIGNMENTS

## APPLICANT: Jochen, Salfeld et al. FITTE OF INVENTION: Human Antibodies That Bi FILE REFERENCE: BBI-093CP CURRENT APPLICATION NUMBER: US/09/534,717 CURRENT FILING DATE: 2000-03-24 EARLIER APPLICATION UNMBER: 60/126,603 EARLIER FILING DATE: March 25, 1999 NUMBER OF SEQ ID NOS: 675 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 494 LENGTH: 13 ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-534-717-494 Sequence 135, Application US/08465325 Patent No. 5686563 Query Match Best Local Similarity GENERAL INFORMATION: Sequence 494, Application US/09534717 Patent No. 6914128 7 IGSNYVK 13 5 LGEEYVK 11 Conservative 35.3%; Score 24; DB 2; Le Pred. No. 8.3e+02; 1; Mismatches 2; Bind Human IL-12 And Methods For Producing Length 13; Indels 0 Gaps 0

Henderson, Farabow, N.W. Suite 700

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RESULT 3
US-09-115-737-135
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 Sequence 135, Application US/09115737

Sequence 136, Application US/09115737

Patent No. 6348445

GENERAL INFORMATION:
APPLICANT: U. Prasad Kari
Taffy J. Williams
Michael McLane
Michael McLane
Tiffle OF INVENTION: Biologically Active Peptides With Reduced
Toxicity in Animals and a Method for Preparing Same
 Matches
 Query Match
 CLASSIFICATION: 514

PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/184,462

PILING DATE: 18-JAN-94

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/891,201

PILING DATE: 01-JUN-92

ATTORNEY/AGENT INFORMATION:
 SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
TYPE: amino acid
STRANUEDNESS: single
TOPOLOGY: linear
 TELEFAX: (202) 408-4400 INFORMATION FOR SEQ ID NO: 135:
 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,325
FILING DATE: 05-JUN-1995
 MOLECULE TYPE:
 NAME: Fordis, Jean B
REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATION GATA:
SOFTWARE: Patentin Release #1.0, Version #1.3
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/115,737
FILING DATE: 15-Jul-1998
CLASSIFICATION STATE: 15-Jul-1998
CLASSIFICATION NUMBER: 08/465,330
FILING DATE: 05-JUN-1995
APPLICATION NUMBER: 08/184,462
FILING DATE: 18-JAN-94
APPLICATION NUMBER: 08/184,462
FILING DATE: 18-JAN-94
APPLICATION NUMBER: 08/184,462
 Local Similarity
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 CORRESPONDENCE ADDRESS:
 NUMBER OF SEQUENCES: 156
 3 KYLGEEYVK 11
 COUNTRY: USA
ZIP: 20005-3315
 ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P.
STREET: 1300 I Street, N.W. Suite 700
 STATE: D.C
 CITY: Washington
APPLICATION NUMBER: 07/891,201
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 peptide
 33.8%;
 3; Mismatches
 Score 23; DB 1; Length 13; Pred. No. 1.2e+03;
 05387.0021-03000
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Sequence 7, Application US/09759484
Patent No. 6858579
GENERAL INFORMATION:
APPLICANT: William Harvey Research Limited
TITLE OF INVENTION: Anti-inflammatory compounds
FILE REFERENCE: P019602W0
CURRENT APPLICATION NUMBER: US/09/759,484
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 7
 Query Match
Best Local Similarity
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 ; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 135:
US-09-115-737-135
 US-09-759-484-7
 RESULT 5
 US-09-060-450-8
 US-09-060-450-8
 APPLICANT: Nardin, Elizabeth
APPLICANT: Moreno, Alberto
TITLE OF INVENTION: UNIVERSAL T-CELL EPITOPES FOR ANTI-MALARIAL VACCINES
FILE REFERENCE: 5986/18615-US1
CURRENT APPLICATION NUMBER: US/09/060,450
CURRENT FILING DATE: 1998-01-21
EARLIER APPLICATION NUMBER: 60/033,916
EARLIER FILING DATE: 1997-01-21
NUMBER OF SEQ ID NOS: 11
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO B
LENGTH: 13
TYPE: PRT
 Sequence 8, Application US/09060450 Patent No. 6669945
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 GENERAL INFORMATION:
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 FEATURE: OTHER INFORMATION: UD4 peptide containing DR 4 allele specific binding OTHER INFORMATION: motifs for use as indicator peptide.
 ORGANISM: Artificial Sequence
 TELEFAX: (202) 408-4400 INFORMATION FOR SEQ ID NO: 135:
 Local Similarity nes 4; Conserv
 RILING DATE: 01-JUN-92
ATTORNEY/AGENT INFORMATION:
NAME: Fordis, Jean B
REGISTRATION NUMBER: 32,984
REFERENCE/DOCKET NUMBER: 05387.0021-06000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408.4000
 SEQUENCE CHARACTERISTICS:
 1 YEKYLGEEYVKA 12
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 LENGTH: 13 amino acids
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STRANDEDNESS: single
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 KKFGKKFVK 9
 TOPOLOGY: linear
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Pred. No. 1
 Score 23; DB 2;
Pred. No. 1.2e+03;
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; APPLICANT: KLIEN, MICHEL A.;BOUX, HEATHER A.;COCKLE,
;STEPHEN A.;LOOSMORE, SHEENA M.;ZEALEY, GAVIN R.
;STEPHEN A.;LOOSMORE, SHEENA M.;ZEALEY, GAVIN R.
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 ; ORGANISM: Escherichia coli
US-10-146-305-14
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US-10-146-305-14
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 US-09-759-484-7
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Best Local S
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APPLICANT: YUHAN CORPORATION
 Sequence 14, Application US/10146305
Patent No. 6939956
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FILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 275,376
FILING DATE: 23-NOV-1988
 CURRENT APPLICATION NUMBER: US/10/146,305
CURRENT FILING DATE: 2002-05-15
PRIOR APPLICATION NUMBER: KR 10-2001-26634
PRIOR FILING DATE: 2001-05-16
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 TITLE OF INVENTION: A VARIABLE REGION OF THE MONOCLONAL ANTIBODY AGAINST THE HBV
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 7 QEYVQTV 13
 7 EEYVKAV 13
 1 YEKYL 5
 4; Conservative
 4; Conservative
 Similarity
YESYL 12
 33.8%;
 33.8%;
 33.8%;
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 1; Mismatches
 Score 23; DB 2;
Pred. No. 1.2e+03;
 Score 23; DB 7;
Pred. No. 1.2e+03
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 Mismatches
 Mismatches
 1; Indels
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 Length 13;
 Indels
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 0
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 Gaps
 Gaps
 Gaps
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 0
 0
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US-08-596-387B-22
 RESULT 8
US-08-480-190-40
 RESULT 9
 US-08-480-190-40
 Best Loc
Matches
 Sequence 22, Application US/08596387B Patent No. 5869270
 Sequence 40, Application US/08480190 Patent No. 5827516
 Query Match
 ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,190
 FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/077,255
FILING DATE: June 15, 1993
APPLICATION NUMBER: 07/925,460
FILING DATE: AUGUST 11, 1992
ATTORNEY/AGENT INFORMATION:
 GENERAL INFORMATION:
 TELEX: 200154
INFORMATION FOR SEQ ID NO:
 GENERAL INFORMATION:
 APPLICANT: Wong, Hing
TITLE OF INVENTION: MHC COMPLEXES AND USES THEREOF
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
 APPLICANT: Rhode, Peter R. APPLICANT: Jiao, Jin-An APPLICANT: Burkhardt, Martin
 SEQUENCE CHARACTERISTICS:
 NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00246/168001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
 APPLICANT: Jack L. Strominger
TITLE OF INVENTION: IMMUNOMODULATORY PEPTIDES
NUMBER OF SEQUENCES: 274
CORRESPONDENCE ADDRESS:
ADDRESSEB: Fish & Richardson
 APPLICANT:
APPLICANT:
 APPLICANT:
 ADDRESSEE:
 STRANDEDNESS:
TOPOLOGY: li
 STREET: 225
CITY: Boston
 COUNTRY: U.S.A
 Local
 ENGTH:
 Similarity 42.9
3; Conservative
 amino acid
 EQYLKKI 10
 EEYVKAV 13
 Massachusetts
 225 Franklin Street
 Roman M. Chicz
Dario A. A. Vignali
Mary L. Hedley
 Lawrence J. Stern
 Robert G. Urban
 linear
Dade International, Inc
 32.4%;
 40:
 Score 22; I
Pred. No. 1.
 Mismatches
 5.
0)
 DB 1; I
1.9e+03;
 Length 13;
 Indels
 0
 Gaps
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STREET:
CITY: De
STATE: I
COUNTRY:

Deerfield : Illinois

1717 Deerfield Road

60015

USA

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US-08-488-379-40; Sequence 40, Application US/08488379
 RESULT 10
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TOPOLOGY:
US-08-596-387B-22
 S
 TELEPHONE: (708) 267-5300
TELEFAX: (708) 267-5376
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
 GENERAL INFORMATION:
APPLICANT: Robert
APPLICANT: Roman M
APPLICANT: Dario A
APPLICANT: Mary L.
 Best Local Similarity
Matches 4; Conserv
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/596,387B
FILING DATE:
CLASSIFICATION UNMBER: PCT/US95/09816
FILING DATE: 31-JUL-1995
PRIOR APPLICATION NUMBER: US 08/382,454
FILING DATE: 01-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/382,454
FILING DATE: 01-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/283,302
FILING DATE: 02-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: PEARSON, LOUISE S.
REGISTRATION NUMBER: 32,369
REFERENCE/DOCKET NUMBER: 32,369
REFERENCE/DOCKET NUMBER: 37R-4665-CIP2
TELEPHONE: (708) 267-5300
TELEPHONE: (708) 267-5300
 Patent No.
 Query Match
 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
 APPLICANT: Jack L. Strominger
TITLE OF INVENTION: IMMUNOMODULATORY PEPTIDES
NUMBER OF SEQUENCES: 274
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 APPLICANT:
 CITY: Boston
 STREET:
 COUNTRY:
 E: Massachusetts
TRY: U.S.A.
02110-2804
 v
 KYLG 6
 KYLG 8
 225 Franklin Street
 Mary L. Hedley
Lawrence J. Stern
 Robert G. Urban
 Roman M. Chicz
Dario A. A. Vignali
 32.4%; Score 22; DB ilarity 100.0%; Pred. No. 1.9 Conservative 0; Mismatches
 unknown
 Version #1.30
 DB 1; Le
 Length 13;
 Indels
 0
 Gaps
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RESULT 12 US-09-067-615-22

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 ; OTHER INFORMATION: Description of Artificial Sequence: humanized
US-09-171-945-33
 US-09-171-945-33
 RESULT 11
 US-08-488-379-40
 NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn Ve
SEQ ID NO 33
 GENERAL INFORMATION:
 Sequence 33, Application US/09171945 Patent No. 6277599
 Query Match 32.4%;
Best Local Similarity 42.9%;
Matches 3; Conservative
 Query Match
Best Local 9
 APPLICANT: Emery, Stephen
APPLICANT: Copley, Clive Graham
APPLICANT: Copley, Clive Graham
APPLICANT: Copley, Clive Graham
APPLICANT: Edge, Michael Derek
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said
TITLE OF INVENTION: Monoclonal Antibody to CEA
CURRENT APPLICATION NUMBER: US/09/171,945
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: GB9703103.3
PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB9609405.7
 FILING DATE: AUGUST 11, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00246/168001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-8906
TELEPHONE: (617) 542-8906
 PRIOR FILING DATE: 1996-05-04
PRIOR APPLICATION NUMBER: PCT/GB97/01165
 PRIOR FILING DATE:
 TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
 INFORMATION FOR SEQ ID NO:
 LENGIH: 13
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/077,255
FILING DATE: June 15, 1993
APPLICATION NUMBER: 07/925,460
 SEQUENCE CHARACTERISTICS:
 TOPOLOGY:
 STRANDEDNESS
 APPLICATION NUMBER:
 CLASSIFICATION:
 LENGTH:
 1 YEKYLGEEY 9
 Similarity 44.
4; Conservative
 7 EEYVKAV 13
|:|:| :
1 EQYLKKI 10
 amino acid
 YAGYLAMDY 13
 PatentIn Ver.
 linear
 1997-04-29
 32.4%;
 US/08/488,379
 40:
 Score 22; DB Pred. No. 1.9e 3; Mismatches
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 Score 22; DB 2;
Pred. No. 1.9e+03;
 Mismatches
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1.9e+03;
 DB 2; Length 13,
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 Indels
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Application US/09067615

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RESULT 13
US-09-374-671A-108
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 US-09-067-615-22
 Sequence 108, Application US/09374671A Patent No. 6489118
GENERAL INFORMATION:
 Sequence 22,
Patent No. 63
 Matches
 Query Match
 TELEFAX: (708) 267-5376
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
 GENERAL INFORMATION:
 TTORNEY, LOUISE 2,369
NAME: PEATSON, LOUISE 32,369
REGISTRATION NUMBER: STR-4665-CIP2
REFERENCE/DOCKET NUMBER: STR-4665-CIP2
TELECOMMUNICATION INFORMATION:
709000E: (708) 267-5376
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/382,454
FILING DATE: 01-FEB-1995
PRIOR APPLICATION DATA:
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
 FILING DATE: 29-JUL-1994
ATTORNEY/AGENT INFORMATION:
 NUMBER OF SEQUENCES: 1:
 APPLICANT:
 APPLICANT:
 STREET: 1.1.
CTTY: Deerfield
TTY: Trinois
 / Match 32.4%; Score 22; Local Similarity 100.0%; Pred. No. 1.9
 LENGTH: 13 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
 APPLICATION NUMBER: US/09/067,615 FILING DATE:
 APPLICATION NUMBER: US 08/283,302 FILING DATE: 29-JUL-1994
 FILING DATE:
 COUNTRY: USA
 CLASSIFICATION:
 ADDRESSEE:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Amy E. Mandragouras
STREET: 28 State Street
 APPLICANT: Morgenstern, Jay P.
 NUMBER OF SEQUENCES:
 TITLE OF
 60015
 INVENTION: M
 E: Dade International, Inc
1717 Deerfield Road
 Rhode, Peter R.
Jiao, Jin-An
Burkhardt, Martin
 INVENTION: Allergenic Protein and Peptides from Dog
Dander and Uses Therefor
 Bizindaukas, Christine B. Brauer, Andrew W.
 Konieczny, Andrzej
 MHC COMPLEXES AND USES THEREOF
 08/596,387
 109
 DB 2;
 1.9e+03;
 0; Indels
 Length 13;
 0;
 Gaps
 0
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 Ś
 US-08-475-399A-40
 US-09-374-671A-108
 Matches
 Patent No.
 Sequence 40, Application US/08475399A
 Query Match
 GENERAL INFORMATION:
 INFORMATION FOR SEQ ID NO: 108:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
TYPE: amino acid
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,399A
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/077,255
 COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
 NUMBER OF SEQUENCES: 276
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
 APPLICANT: Urban, Robert G.
APPLICANT: Chicz, Roman M.
APPLICANT: Vignali Dario A.A
APPLICANT: Hedley, Mary L.
APPLICANT: Stern, Lawrence J.
 APPLICANT: Strominger, Jack L.
TITLE OF INVENTION: IMMUNOMODULATORY PEPTIDES
 APPLICATION NUMBER: US/09/374,671A
FILING DATE: 16-Aug-1999
PRIOR APPLICATION NUMBER: US 08/156,549
FILING DATE: 1993-NOV-22
APPLICATION NUMBER: US 07/999,712
FILING DATE: 1992-DEC-31
ATTORNEY/AGENT INFORMATION:
NAME: DIGIOTGIO, Jeanne M.
REGISTRATION NUMBER: 41,710
REFERENCE/DOCKET NUMBER: IMI-026C2CNCPA
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 742-4214
TELEPAX: (617) 742-4214
 STREET: 225
CITY: Boston
 Match 32.4%;
Local Similarity 50.0%;
 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: ASCII Text

CURRENT APPLICATION DATA:
 MOLECULE TYPE: peptide FRAGMENT TYPE: internal
 SEQUENCE DESCRIPTION: SEQ ID NO: 108:
 6 GEEYVKAV 13
 650903
 4.
 CITY: Boston
STATE: MA
COUNTRY: USA
 GKWYLKAM 8
 M
 TOPOLOGY: linear
 ZIP: 02109
 225 Franklin Street
 Conservative
 Dario A.A.
 Mismatches
 Score 22; DB 2;
Pred. No. 1.9e+03;
 1; Indels
 Length 13;
 0
 Gaps
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REFERENCE/DOCKET NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00246/168003
FELECOMMUNICATION INFORMATION:
FELEPHONE: 617/542-890
FELEX: 200154
FILEX: 200154
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
TYPE: amino acid
TYPE: amino acid
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TYPE: amino acid
 APPLICANT: PYKETT, Mark J.

APPLICANT: PYKETT, Mark J.

APPLICANT: SCADDEN, David T.

APPLICANT: POZNANSKY, Mark C.

ITITLE OF INVENTION: LYMPHOID TISSUE-SPECIFIC CELL PRODUCTION

ITITLE OF INVENTION: FROM HEMATOPOIETIC PROGENITOR CELLS IN THREE-DIMENSIONAL

ITITLE OF INVENTION: FROM HEMATOPOIETIC PROGENITOR CELLS IN THREE-DIMENSIONAL

ITITLE OF INVENTION NUMBER: US5/99/574,749B

CURRENT FILING DATE: 2002-05-31

PRIOR APPLICATION NUMBER: US 60/107,972

PRIOR APPLICATION NUMBER: US 60/107,972

PRIOR APPLICATION NUMBER: US 60/107,972

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR FILING DATE: 1999-11-12

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR APPLICATION NUMBER: US 09/524,749

PRIOR FILING DATE: 1999-11-12

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 45

SEQ ID NO 45

LENGTH: 13

TYPE: PRIOR APPLICATION SECOND NOWN VERSION 3.0
Search completed: June 13, 2006, 03:12:35 Job time : 51 secs
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 ફ
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US-09-574-749B-45
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 Sequence 45, Application US/09574749B Patent No. 6548299
 Query Match 32.4%; Score 22; DB 2; Length 13; Best Local Similarity 42.9%; Pred. No. 1.9e+03; Matches 3; Conservative 3; Mismatches 1; Indels
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Best Local Similarity
 GENERAL INFORMATION:
 ORGANISM: Artificial Sequence FEATURE:
 FILING DATE: 15-JUN-1993
APPLICATION NUMBER: 07/925,460
FILING DATE: 11-AUG-1992
ATTORNEY/AGENT INFORMATION:
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 7 EEYVK 11
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80.0%; Pred. No. 1.9e+03;
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Result
 Minimum DB
Maximum DB
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 Total number of hits satisfying chosen parameters:
 OM protein -
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 Scoring table:
 Perfect score:
 Title:
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 9
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Score
 seq length:
seq length:
 protein search, using sw
 Query
Match
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 US-10-612-162A-4
68
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4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
 Minimum Match 0%
Maximum Match 100%
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 GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd
 Length
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 US-10-612-162-4
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US-11-105-268-3
US-10-280-139-11
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US-10-930-300-113
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US-10-936-647-889
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Appl
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| 23               | 23                 | 23                | 23                | 23                | 23              | 23                 | 23                | 23                | 23                | 23                | 23                | 23                | 23               | 23               | 23                | 23               | 23               |
| •                | •                  | •                 |                   | 33.8              | •               | 33.8               |                   | •                 | 33.8              |                   | 33.8              |                   |                  | •                | 33.8              | •                | 33.8             |
| 13               | 13                 | 13                | 13                | 13                | 13              | 13                 | 13                | 13                | 13                | 13                | 13                | 13                | 13               | 13               | 13                | 13               | 13               |
| ហ                | ഗ                  | თ                 | ທ                 | ហ                 | S               | S                  | u                 | 5                 | ហ                 | ഗ                 | S                 | ហ                 | v                | v                | 4                 | 4.               | 4.               |
| US-10-716-189-8  | US-10-948-707-1360 | US-10-948-707-911 | US-10-948-707-745 | US-10-958-216-716 | US-10-296-313-1 | US-10-470-048B-603 | US-10-930-300-112 | US-10-930-300-109 | US-10-930-300-107 | US-10-930-300-105 | US-10-930-300-102 | US-10-930-300-100 | US-10-930-300-98 | US-10-834-666-44 | US-10-469-837-14  | US-10-469-837-13 | US-10-469-837-12 |
| Sequence 8, Appl | Sequence 1360,     | Sequence 911, A   | 745,              | Sequence 716, A   | Sequence 1, App | Sequence 603, A    | Sequence 112, A   | Sequence 109, A   | Sequence 107, A   | Sequence 105, A   | Sequence 102, A   | Sequence 100, A   | Sequence 98, Ap  | Sequence 44, Ap  | Sequence 14, Appl | Sequence 13, Ap  | Sequence 12, Ap  |
| 1 i              | ď                  | ďď                | App               | ďď                | 1               | ďď                 | ďď                | ďď                | ď                 | ďď                | ďď                | ďď                | ģ                | Ď                | Ď                 | Ď.               | Ţġ               |

ALIGNMENTS

## Sequence 3, Application US/10823253 Publication No. US20050002934A1 GENERAL INFORMATION: APPLICANT: Reed, Jennifer TITLE OF INVENTION: RECOMBINANT IL-9 ANTIBODIES AND USES THEREOF FILE REFERENCE: 10271-112-999 CURRENT APPLICATION NUMBER: US/10/823,253 CURRENT FILING DATE: 2004-04-12 PRIOR APPLICATION NUMBER: 60/462,259 PRIOR FILING DATE: 2003-04-11 PRIOR APPLICATION NUMBER: 60/477,797 PRIOR APPLICATION NUMBER: 60/477,797 PRIOR APPLICATION NUMBER: 60/477,797 PRIOR APPLICATION NUMBER: 60/477,797 PRIOR FILING DATE: 2003-06-10 NUMBER OF SEQ ID NOS: 60 SOFTWARE: Patentin version 3.2 SEQ ID NO 3 밁 ફ US-10-823-253-3 US-10-612-162-4 RESULT 1 US-10-612-162-4 PRIOR APPLICATION NUMBER: 10230550.1 PRIOR FILING DATE: 2002-07-05 PRIOR APPLICATION NUMBER: 03011334.4 PRIOR FILING DATE: 2003-05-19 NUMBER OF SEQ ID NOS: 4 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 4 Sequence 4, Application US/10612162 Publication No. US20040014145A1 GENERAL INFORMATION: APPLICANT: Dade Behring Marburg GmbH TITLE OF INVENTION: antibodies, their preparation and use FILE REFERENCE: 2002/B001 Matches Best Query Match CURRENT APPLICATION NUMBER: US/10/612,162 CURRENT FILING DATE: 2003-07-03 LENGTH: 13 TYPE: PRT ORGANISM: Homo sapiens Local 1 YEKYLGEEYVKAV 13 Similarity YEKYLGEEYVKAV 13 Conservative 100.0%; Score 68; DB 4; 100.0%; Pred. No. 0.0002; <u>,</u> Mismatches 0; Length 13; Indels 0; Gaps

0

LENGTH: 13

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CURRENT APPLICATION NUMBER: US/10/823,810
CURRENT FILING DATE: 2004-04-12
PRIOR APPLICATION NUMBER: 60/462,307
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 60/477,801
PRIOR FILING DATE: 2003-06-10
NUMBER OF SEQ ID NOS: 63
SOPTWARE: Patentin version 3.2
SOPTWARE: Patentin version 3.2
LENGTH: 13
LENGTH: 13
TYPE: PRT
밁
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 ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-823-253-3
 ; ORGANISM: Homo sapiens US-10-823-810-3
 US-10-823-810-3
 밁
 US-11-105-268-3
 CURRENT APPLICATION UNBER: US/11/105,268
CURRENT FILING DATE: 2005-04-12
PRIOR APPLICATION UNMBER: 60/561,845
PRIOR FILING DATE: 2004-04-12
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3.
 Sequence 3, Application US/11105268
Publication No. US20050260204A1
GENERAL INFORMATION:
APPLICANT: Allan, Christian
TITLE OF INVENTION: ANTI-IL-9 ANTIBODY FORMULATIONS
FILE REFERENCE: 10271-126-999
 Sequence 3, Application US/10823810 Publication No. US20050147607A1 GENERAL INFORMATION:
 Query Match 44.1%;
Best Local Similarity 62.5%;
Matches 5; Conservative
 Query Match
Best Local :
 Query Match 44.1%;
Best Local Similarity 62.5%;
Matches 5; Conservative
 Matches
 APPLICANT: Reed, Jennifer
TITLE OF INVENTION: METHODS OF PREVENTING OR TREATING RESPIRATORY CONDITIONS
FILE REFERENCE: 10271-113-999
 LENGTH: 13
TYPE: PRT
ORGANISM: Homo sapiens
 4 YLGEEYVK 11
 4 YLGEEYVK 11
 Similarity 62.5%;
5; Conservative
 4 YLGEEYVK 11
 YYGSDYVK 10
YYGSDYVK 10
 YYGSDYVK 10
 Score 30; DB 6;
Pred. No. 3.8e+02;
1; Mismatches 2
 Score 30; DB 5; Lo
Pred. No. 3.8e+02;
1; Mismatches 2;
 Score 30; DB 5;
Pred. No. 3.8e+02;
1; Mismatches 2
 Length 13;
 Length 13
 Length 13;
 Indels
 Indels
 AND USES THEREOF
 Indels
 0
 0;
 0
 Gaps
 Gaps
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 0
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 US-10-930-300-101
 RESULT 7
 US-10-823-253-12
 US-10-823-253-12
 US-10-280-139-11
 US-10-280-139-11
 Query Match
Best Local S
Matches 4
 Sequence 12, Application US/10823253
Publication No. US20050002934A1
GENERAL INFORMATION:
APPLICANT: Reed, Jennifer
TITLE OF INVENTION: RECOMBINANT IL-9 ANTIBODIES AND USES THEREOF
FILE REFERENCE: 10271-112-99
CURRENT APPLICATION NUMBER: US/10/823,253
CURRENT FILING DATE: 2004-04-12
PRIOR APPLICATION NUMBER: 60/462,259
PRIOR FILING DATE: 2003-04-11
 Sequence 11, Application US/10280139
Publication No. US20040081963A1
GENERAL INFORMATION:
APPLICANT: Wang, Yan
APPLICANT: Wang, Yan
APPLICANT: WJ Bioworks Incorporated
TITLE OF INVENTION: Improved Ssc7-Polymerase Conjugate Proteins
FILE REPERENCE: 020130-001200US
FILE REPERENCE: 020130-001200US
CURRENT APPLICATION NUMBER: US/10/280,139
CURRENT FILING DATE: 2002-10-23
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 11
 Sequence 101, Application US/10930300 Publication No. US20050014138A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 12
 Query Match
Best Local Similarity
Matches 5; Conserv
 PRIOR APPLICATION NUMBER: 60/477,797
PRIOR FILING DATE: 2003-06-10
NUMBER OF SEQ ID NOS: 60
 LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence:S807d OTHER INFORMATION: alignment consensus peptide
 ORGANISM: Homo sapiens
 TYPE: PRT
 LENGTH:
 CORRESPONDENCE ADDRESS
 NUMBER OF SEQUENCES:
 APPLICANT: Rath, Matthias
TITLE OF INVENTION: METHOD OF PRODUCING VACCINES FROM PROTEIN SIGNAL
 3 KYLGEE 8
 4 YLGEEYVK 11
 4 KYKGEE
 Similarity 4; Conserv
 YYGSDHVK 10
 Conservative
 Conservative
INHOUSE IP MANAGEMENT
 38.2%;
 36.8%;
 OLIGOPEPTIDES
 0; Mismatches
 Score 25; DB 5;
Pred. No. 2.6e+03;
 Score 26; DB 4;
Pred. No. 1.8e+03;
 Mismatches
 Length 13;
 Length 13
 Indels
 and Sac7e
 0
 0
 Gaps
 Gaps
 0
 0
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CITY: Palo Alto

STREET: 280 Colorado Avenue

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RESULT 8
US-10-930-300-108
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 US-10-930-300-101
 Sequence 108, Application US/10930300 Publication No. US20050014138A1 GENERAL INFORMATION:
 Matches
 Query Match
 TELEFAX: 650-322-7389 INFORMATION FOR SEQ ID NO: 101:
 Match 36.8%;
Local Similarity 45.5%;
es 5; Conservative
 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy Disk, 3.50 inch, 1.44MB Storage COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: WINDOWS 95

SOFTWARE: MS WORD 6.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/930,300

FILING DATE: 30-Aug-2004

CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,248
FILING DATE: 14-JAN-1994
ATTORNEY/AGENT INFORMATION:
NAME: ALI KAMAREI
 NUMBER OF SEQUENCES: 360
CORRESPONDENCE ADDRESS:
ADDRESSE: INHOUSE IP MANAGEMENT
STREET: 280 Colorado Avenue
 STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 101:
 NAME: ALI KAMAREI
REGISTRATION NUMBER: 37000
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-322-7371
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,248
FILING DATE: 14-JAN-1994
ATTORNEY/AGENT INFORMATION:
 APPLICANT: Rath, Matthias
TITLE OF INVENTION: METHOD OF PRODUCING VACCINES FROM PROTEIN SIGNAL
OLIGOPEPTIDES
 CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM: Disk, 3.50 inch, 1.44MB Storage
 SEQUENCE CHARACTERISTICS:
 1 YEKYLGEEYVK 11
 STREET: 280 Colore
CITY: Palo Alto
STATE: California
 APPLICATION NUMBER: US/10/930,300 FILING DATE: 30-Aug-2004 CLASSIFICATION: <Unknown>
 COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: WINDOWS 95 SOFTWARE: MS WORD 6.0
REGISTRATION NUMBER: 37000
 COUNTRY: USA
 TYPE: amino acid
 ZIP: 94301
 COUNTRY: USA
 STATE: California
 LENGTH: 13
 0; Mismatches
 Score 25; DB 5;
Pred. No. 2.6e+03;
 10262-1US
 6; Indels
 Length 13;
 Gaps
 0
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 ; SEQUENCE DESCRIPTION: SEQ ID NO: 113: US-10-930-300-113
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 RESULT 9
US-10-930-300-113
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 US-10-930-300-108
 Matches
 Sequence 113, Application US/10930300
Publication No. US20050014138A1
GENERAL INFORMATION:
APPLICANT: Rath, Matthias
 Query Match
Best Local Similarity
 Matches
 Query Match
 TELEFAX: 650-322-7389
INFORMATION FOR SEQ ID NO: 113:
 INFORMATION FOR SEQ ID NO: 108: SEQUENCE CHARACTERISTICS:
 PRIOR APPLICATION DATA:

APPLICATION UNMBER: 08/182,248

FILING DATE: 14-JAN-1994

ATTORNEY/AGENT INFORMATION:

NAME: ALI KAMAREI

REGISTRATION NUMBER: 37000

REGISTRATION NUMBER: 10262-11

TELEPRENCE/DOCKET NUMBER: 10262-11

TELEPHONE: 650-322-7371
 36.8%;
Local Similarity 45.5%;
es 5; Conservation
 COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: MS WORD 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/930,300
FILING DATE: 30-Aug-2004
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk, 3.50 inch, 1.44MB Storage
 SEQUENCE CHARACTERISTICS:
 CORRESPONDENCE ADDRESS
 NUMBER OF SEQUENCES:
 TITLE OF INVENTION: METHOD OF PRODUCING VACCINES FROM PROTEIN SIGNAL
 SEQUENCE DESCRIPTION: SEQ ID NO: 108:
 TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-322-7371
 N
 1 YEKYLGEEYVK 11
 YEKYLGEEYVK 11
 STREET: 280 Colorado Avenue
CITY: Palo Alto
STATE: California
YSKYLDNRRTK 12
 STRANDEDNESS: single TOPOLOGY: linear
 TYPE: amino acid
 YSKYLDNRRAK 12
 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
 CLASSIFICATION: <Unknown>
 COUNTRY: USA
 TELEFAX: 650-322-7389
 LENGTH: 13
 ZIP: 9430:
 ADDRESSEE: INHOUSE IP MANAGEMENT
 REFERENCE/DOCKET NUMBER: 10262-1US
 LENGTH: 13
 Conservative
 36.8%;
 OLIGOPEPTIDES
 0; Mismatches
 Score 25; DB 5;
Pred. No. 2.6e+03;
 Score 25; DB 5;
Pred. No. 2.6e+03;
 Mismatches
 10262-1US
 Length 13
 6; Indels
 Length 13;
 0,
 0
 Gaps
 Gaps
 0
 0,
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 US-11-105-268-12
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 ; ORGANISM: Escherichia coli
US-10-946-647-889
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 US-10-823-810-12
 RESULT 12
 US-10-946-647-889
 US-10-823-810-12
 Publication No. US20050186217A1
GENERAL INFORMATION:
APPLICANT: EMERY, DARYLL A.
APPLICANT: STRAUB, DARREN E.
APPLICANT: WONDERLING, LAURA
TITLE OF INVENTION: COMPOSITIONS PRODUCED USING ENTERIC PATHOGENS AND METHODS OF USE
FILE REFERENCE: 293.00340101
CURRENT APPLICATION NUMBER: US/10/946,647
CURRENT APPLICATION NUMBER: 60/504,119
PRIOR APPLICATION NUMBER: 60/504,119
PRIOR PILING DATE: 2003-09-19
NUMBER OF SEQ ID NOS: 1448
SOFTWARE: Patentin version 3.3
SEQ ID NO 889
LENGTH: 13
 Sequence 12, Application US/11105268
Publication No. US20050260204A1
GENERAL INFORMATION:
APPLICANT: Allan, Christian
TITLE OF INVENTION: ANTI-IL-9 ANTIBODY FORMULATIONS AND USES THEREOF
FILE REFERENCE: 10271-126-999
 APPLICANT: Reed, Jennifer
TITLE OF INVENTION: METHODS OF PREVENTING OR TREATING RESPIRATORY CONDITIONS
FILE REFERENCE: 10271-113-999
CURRENT APPLICATION NUMBER: US/10/823,810
CURRENT FILING DATE: 2004-04-12
PRIOR APPLICATION NUMBER: 60/462,307
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 60/477,801
PRIOR APPLICATION NUMBER: 60/477,801
PRIOR TILING DATE: 2003-06-10
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin version 3.2
SEQ ID NO 12
 Sequence 12, Application US/10823810 Publication No. US20050147607A1 GENERAL INFORMATION:
 Query Match
Best Local
 Query Match
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 Matches
CURRENT APPLICATION NUMBER: US/11/105,268
 LENGTH: 1:
TYPE: PRT
 LENGTH: 13
TYPE: PRT
ORGANISM: Homo sapiens
 Local Similarity 44.4%;
nes 4; Conservarion
 Local Similarity
 w
 4 YLGEEYVKA 12
 YASPEYIQA 9
 YLGEEYVK 11
 Application US/10946647
o. US20050186217A1
 Conservative
 36.8%;
 Score 25; DB 5;
Pred. No. 2.6e+03;
 Score 25; DB 5; Pred. No. 2.6e+03; 2; Mismatches 2
 Length 13;
 Length 13;
 Indels
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 Gaps
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Sequence 302, Application US/11153143A
Publication No. US20060051396A1
GENERAL INFORMATION:
APPLICANT: Paul T. Hamilton
APPLICANT: Mark W. Grinstaff
APPLICANT: Daniel J. Kenan
APPLICANT: Dale J. Christensen
TITLE OF INVENTION: Biofunctional Coatings
FILE REFERENCE: AFF005
 PRIOR APPLICATION NUMBER: 60/561,845
PRIOR FILING DATE: 2004-04-12
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn version 3.2
SEQ ID NO 12
LENGTH: 13
TYPE: PRT
ORGANISM: Homo sapiens
 RESULT 13
US-11-152-974A-302
 Query Match
Best Local Similarity
Watches 4; Conserva
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 S
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 5
 US-11-152-974A-302
 US-11-105-268-12
 US-11-153-143A-302
 NUMBER OF SEQ ID NOS: 558
SOFTWARE: FRACTSEQ for Windows Version 4.0
SEQ ID NO 302
LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Sequence 302, Application US/11 Publication No. US20060051395A1
 APPLICANT: Wayne F. Beyer, Jr.
APPLICANT: Robin Hyde-DeRuyscher
APPLICANT: Paul T. Hamilton
APPLICANT: Ray Edward Benson
TITLE OF INVENTION: IFMBS to Promote the Specific Attachment of Target Analytes to the
TITLE OF INVENTION: of Orthopedic Implants
FILE REFERENCE: APFO06
CURRENT APPLICATION NUMBER: US/11/152,974A
CURRENT FILING DATE: 2005-06-15
PRIOR APPLICATION NUMBER: 60/580,019
PRIOR APPLICATION NUMBER: 60/581,338
PRIOR FILING DATE: 2004-06-16
PRIOR PILING DATE: 2005-02-09
PRIOR FILING DATE: 2005-02-09
PRIOR FILING DATE: 2005-02-09
PRIOR FILING DATE: 2005-02-09
PRIOR FILING DATE: 2005-02-09
 PRIOR APPLICATION NUMBER: 60/651,747 PRIOR FILING DATE: 2005-02-10
 CURRENT APPLICATION NUMBER: US/11/153,143A CURRENT FILING DATE: 2005-06-15
 CURRENT FILING DATE:
PRIOR APPLICATION NUMBER: 60/580,019
 OTHER INFORMATION: isolated from phage display libraries
 FEATURE:
 1 YEKYLG
 4 YLGEEYVK 11
 4; Conservative
 YDKYSG
 YYGSDHVK 10
 Application US/11152974A
 Conservative
 36.8%;
 2005-04-12
 36.8%;
 Mismatches
 Score 25; DB 6;
Pred. No. 2.6e+03;
 Score 25; DB 6;
Pred. No. 2.6e+03
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 Length 13
 Length 13
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 Gaps
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Sequence 18, Application US/10080980

Publication No. US20030036115A1

GENERAL INFORMATION:

APPLICANT: Bristol-Myers Squibb Company

TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN POTASSIUM CHANNEL BETA-SUBU

TITLE OF INVENTION: K+betaM6, EXPRESSED HIGHLY IN THE SMALL INTESTINE

FILE REFERENCE: D0121 NP

CURRENT APPLICATION NUMBER: US/10/080,980

CURRENT FILING DATE: 2002-02-21

PRIOR APPLICATION NUMBER: US 60/270,132

PRIOR APPLICATION NUMBER: US 60/270,132

PRIOR APPLICATION NUMBER: US 60/278,953

PRIOR FILING DATE: 2001-03-27

NUMBER OF SEQ ID NOS: 74

SOFTWARE: Patentin version 3.0

SEQ ID NO 18

LENGTH: 13

TYPE: PRT

ORGANISM: homo sapiens
 PRIOR FILING DATE: 2004-06-16
PRIOR APPLICATION NUMBER: 60/651,338
PRIOR FILING DATE: 2005-02-09
PRIOR APPLICATION NUMBER: 60/651,747
PRIOR APPLICATION NUMBER: 60/651,747
PRIOR FILING DATE: 2005-02-10
NUMBER OF SEQ ID NOS: 558
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 302
LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: isolated from phage display libraries
US-11-153-143A-302
Search completed: June 13, 2006, 03:29:35 Job time : 182 secs
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 RESULT 15
US-10-080-980-18
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 Query Match 35.3%;
Best Local Similarity 40.0%;
Matches 4; Conservative
 Query Match 36.8%;
Best Local Similarity 66.7%;
Matches 4; Conservative
 1 YEKYLG 6
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7 YDKYSG 12
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 EKYLGEEYVK 11
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 Score 25; DB 6; Length 13; Pred. No. 2.6e+03; 1; Mismatches 1; Indels
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Copyright

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 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 s
 Minimum DB
Maximum DB
 Total number of hits satisfying chosen parameters:
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 Sequence:
 Title:
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 Perfect score:
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 Pred. No. score grea and is der
 Score
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 is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.
 Published Applications AA New:*

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3: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

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6: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

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30, Appl
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RESULT 1 US-10-497-088-5

ALIGNMENTS

Sequence 5, Application US/10497088 Publication No. US20060088520A1 GENERAL INFORMATION:

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| 16 23.5 13 7 US-11-054-072-5013 16 23.5 13 7 US-11-054-072-5013 16 23.5 13 7 US-11-054-072-5013 16 23.5 13 7 US-11-054-072-5013 16 23.5 13 7 US-11-054-072-5014 16 23.5 13 7 US-11-054-072-5016 16 23.5 13 7 US-11-054-072-5017 16 23.5 13 7 US-11-054-072-5019 16 23.5 13 7 US-11-054-072-5020 16 23.5 13 7 US-11-054-072-5020 16 23.5 13 7 US-11-054-072-5020 16 23.5 13 7 US-11-054-072-5022 16 23.5 13 7 US-11-054-072-5022 16 23.5 13 7 US-11-054-072-5023 16 23.5 13 7 US-11-054-072-5024 16 23.5 13 7 US-11-054-072-5024 16 23.5 13 7 US-11-054-072-5026 16 23.5 13 7 US-11-054-072-5026 16 23.5 13 7 US-11-054-072-5027 16 23.5 13 7 US-11-054-072-5026 16 23.5 13 7 US-11-054-072-5027 16 23.5 13 7 US-11-054-072-5026 16 23.5 13 7 US-11-054-072-5027 16 23.5 13 7 US-11-054-072-5026 16 23.5 13 7 US-11-054-072-5027 16 23.5 13 7 US-11-054-072-5026                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 45                | 44              | 43                 | 42                 | 41                 | 40                 | 39       | 38                 | 37                 | 36                 | 35<br>5            | υ<br><b>4</b>      | y<br>W | 32                 | 31                 | 30       | 29                 | 28                 | 27                 | 26 |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------|-----------------|--------------------|--------------------|--------------------|--------------------|----------|--------------------|--------------------|--------------------|--------------------|--------------------|--------|--------------------|--------------------|----------|--------------------|--------------------|--------------------|----|
| .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 13 7 08-11 .5 | 16                | 16              | 16                 | 16                 | 16                 | 16                 | 16       | 16                 | 16                 | 16                 | 16                 | 16                 | 16     | 16                 | 16                 | 16       | 16                 | 16                 | 16                 | 16 |
| 7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11<br>7 US-11                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |                   |                 |                    |                    |                    |                    |          |                    |                    |                    |                    |                    |        |                    |                    |          |                    |                    |                    |    |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 13                | 13              | 13                 | 13                 | 13                 | 13                 | 13       | 13                 | 13                 | 13                 | 13                 | 13                 | 13     | 13                 | 13                 | 13       | 13                 | 13                 | 13                 | 13 |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 7                 | 7               | 7                  | 7                  | 7                  | 7                  | 7        | 7                  | 7                  | 7                  | 7                  | 7                  | 7      | 7                  | 7                  | 7        | 7                  | 7                  | 7                  | `  |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | US-11-264-509A-81 | US-11-233-532-2 | US-11-054-072-5027 | US-11-054-072-5026 | US-11-054-072-5025 | US-11-054-072-5024 |          | US-11-054-072-5022 | US-11-054-072-5021 | US-11-054-072-5020 | US-11-054-072-5019 | US-11-054-072-5018 | -11    | US-11-054-072-5016 | US-11-054-072-5015 | 11       | US-11-054-072-5013 | US-11-054-072-5012 | US-11-054-072-5011 | TT |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 81, Appl          | 2, Appli        | 5027, Ap           | •                  | 5025, Ap           | 5024, Ap           | 5023, Ap | 5022, Ap           | 5021, Ap           | 5020, Ap           | 5019, Ap           | •                  | •      | 5016, Ap           | 5015, Ap           | 5014, Ap | 5013, Ag           | •                  | -                  | •  |

## APPLICANT: Crucell Holland B.V. APPLICANT: Germeraad, Wilfred APPLICANT: Legtenberg, Ton APPLICANT: Legtenberg, Ton APPLICANT: Lekkerker, Annemarie N ITILE OF INVENTION: Antigen presenting cell targeting conjugate, an antigen ITILE OF INVENTION: presenting cell contacted with such conjugate, their use for ITILE OF INVENTION: vaccination or as medicament, and methods for their production or ITILE OF INVENTION: generation ITILE OF INVENTION: US/10/497,088 CURRENT APPLICATION NUMBER: US/10/497,088 CURRENT APPLICATION NUMBER: PCT/EP01/14255 PRIOR APPLICATION NUMBER: PCT/EP01/14255 PRIOR FILING DATE: 2001-11-29 PRIOR FILING DATE: 2002-11-29 PRIOR APPLICATION NUMBER: EP01/204997.9 PRIOR FILING DATE: 2001-12-19 NUMBER OF SEQ ID NOS: 21 SOFTWARE: Patentin version 3.2 SEQ ID NO 5 LANGETH: 13 RESULT 2 US-11-259-950-33 밁 ঠ US-10-497-088-5 Matches Query Match Best Local LENGTH: 13 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: MAGE-1.DR13 specific peptide w 3 KYLGEEYVKA 12 32.4%; Similarity 50.0%; 5; Conservative KYRAEPVTKA 12 Score 22; DB 6; Lo Pred. No. 2.9e+02; D; Mismatches 5; Length 13; Indels 0, Gaps 0

Sequence 33, Application US/11259950
Publication No. US20060088524A1
GENERAL INFORMATION:
APPLICANT: Morriseey, James
APPLICANT: Pureza, Vincent

```
APPLICANT: Zhang, Rumin
APPLICANT: Kong, Rong
TITLE OF INVENTION: PEPTIDE VIRAL ENTRY PEPTIDES
FILE REFERENCE: JB06227US01
CURRENT APPLICATION NUMBER: US/11/264,509A
CURRENT FILING DATE: 2005-11-01
PRIOR APPLICATION NUMBER: 60/624,204
PRIOR FILING DATE: 2004-11-02
NUMBER: OF SEQ ID NOS: 141
SOPTWARE: PATENTIN VERSION 3.3
SEQ ID NO 132
LENGTH: 13
TYDE: DET
US-10-538-066-38
; Sequence 38, Application US/10538066
; Publication No. US20060094649A1
 밁
 RESULT 4
 밁
 ; OTHER INFORMATION: HCV peptide US-11-264-509A-132
 US-11-264-509A-132
 ; OTHER INFORMATION: Helix 10.5 US-11-259-950-33
 PRIOR APPLICATION NUMBER: US 60/536,281
PRIOR FILLING DATE: 2004-01-13
PRIOR PELICATION NUMBER: US 10/465,789
PRIOR FILLING DATE: 2003-06-18
PRIOR FILLING DATE: 2003-06-18
PRIOR FILLING DATE: 2001-11-20
PRIOR PELICATION NUMBER: US 60/252,233
PRIOR FILLING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 61/252,233
PRIOR FILLING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 11/033,489
PRIOR FILLING DATE: 2005-01-11
 Sequence 132, Application US/11264509A Publication No. US20060100148A1 GENERAL INFORMATION:
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 SOFTWARE: PatentIn version 3.3
SEQ ID NO 33
LENGTH: 13
TYPE: PRT
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 Query Match
Best Local (
 APPLICANT: Sligar, Stephen G.
TITLE OF INVENTION: Tissue Factor Composit
FILE REFERENCE: 46-04
CURRENT APPLICATION NUMBER: US/11/259,950
CURRENT FILING DATE: 2005-10-27
PRIOR APPLICATION NUMBER: US 60/622,737
PRIOR FILING DATE: 2004-10-27
PRIOR FILING DATE: 2004-10-27
 APPLICANT: Liu, Rong
APPLICANT: Zhang, Ru
 NUMBER OF SEQ ID NOS: 96
 ORGANISM: Artificial FEATURE:
 ORGANISM: Artificial Sequence
 TYPE: PRT
 FEATURE:
 Match 32.4%;
Local Similarity 80.0%;
 v
 8 EYVK 11
 7 EEYVK 11
 4
 Similarity
 EEYTK 8
 Conservative
 Conservative
 Tissue Factor Compositions and Methods
 30.9%; Score 21; DB 7;
100.0%; Pred. No. 4.3e+
tive 0; Mismatches
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 Score 22; DB 7;
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 0
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Sequence 41, Application US/10538066
Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epinmune Inc.
APPLICANT: Epinmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and TITLE OF INVENTION: Peptides and Compositions FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION WIMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
 Sequence 18, Application US/11038754
Publication No. US20060115855A1
GENERAL INFORMATION:
APPLICANT: Goldknopf, Ira L.
APPLICANT: Sheta, Essam A.
APPLICANT: Appel, Stanley H.
APPLICANT: Simpson, Ericka P.
APPLICANT: Yen, Albert A.
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 US-11-038-754-18
 밁
 RESULT 5
US-10-538-066-41
 멍
 US-10-538-066-38
 RESULT 6
 US-10-538-066-41
 NUMBER OF SEQ ID NOS: 767
SOFTWARE: Patentin version 3.2
SEQ ID NO 38
LENGTH: 13
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 41
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 GENERAL INFORMATION:
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 APPLICANT: Epimmune Inc.

TITLE OF INVENTION: HAP-A1, -A2, -A3, -A24, -B7,

TITLE OF INVENTION: Peptides and Compositions

FILE REFERENCE: 2060.015PC06

CURRENT APPLICATION NUMBER: US/10/538,066

CURRENT FILING DATE: 2005-06-09

PRIOR APPLICATION NUMBER: US 60/432,017

PRIOR FILING DATE: 2002-12-10
 LENGTH: 13
TYPE: PRT
ORGANISM: Unknown
FEATURE:
 TITLE OF INVENTION: AN FK506-BINDING PROTEIN 7 RELATED PROTEIN AS A BIOMARKER FOR TITLE OF INVENTION: NEURODEGENERATIVE DISEASE
FILE REFERENCE: PROT-P0012US
 OTHER INFORMATION: PADRE Peptide
 FEATURE:
OTHER INFORMATION: PADRE Peptide
 ORGANISM: Unknown
 TYPE: PRT
 Local Similarity nes 4; Conserv
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 3 KYLGEEYVKA 12
 KYVAAHTLKA 11
 KYVAANTLKA 11
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 Conservative
 29.4%;
 29.4%;
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 Score 20;
Pred. No.
 Score 20;
Pred. No. 6
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 DB 6;
 DB 6;
 Length 13;
 Length 13;
 and -B44 Tumor Associated Antigen
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Sequence 94, Application US/11264509A
PUDIICATION NO. US20060100148A1
GENERAL INFORMATION:
APPLICANT: Liu, Rong
APPLICANT: Chang, Rumin
APPLICANT: Chang, Rong
TITLE OF INVENTION: PEPTIDE VIRAL ENTRY PEPTIDES
FILE REFERENCE: JB66227US01
CURRENT APPLICATION NUMBER: US/11/264,509A
CURRENT ETLING DATE: 2005-11-01
PRIOR APPLICATION NUMBER: 60/624,204
 US-10-614-959-13
; Sequence 13, Applica
; Publication No. US20
; GENERAL INFORMATION:
 밁
 Query Match
Best Local Similarity
Warches 4; Conserva
 RESULT 8
US-11-264-509A-94
 망
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 ; LENGTH: 13
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-038-754-18
 ; ORGANISM: Homo sapiens
US-10-614-959-13
 PRIOR APPLICATION NUMBER: US 60/632,216
PRIOR FILING DATE: 2004-12-01
NUMBER OF SEQ ID NOS: 21
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 18
 APPLICANT: Adams, Camelia W.
APPLICANT: Devaux, Brigitte
APPLICANT: Eaton, Dan L.
APPLICANT: Hass, Philip E.
APPLICANT: Judice, J. Kevin
APPLICANT: Kirchhofer, Daniel
APPLICANT: Suggett, Shelley
 SEQ ID NO 13
 Matches
 Query Match
 APPLICANT: Kirchhofer, Daniel
APPLICANT: Suggett, Shelley
IIILE OF INVENTION: Human Anti-Factor IX/IXa Antibodies
FILE REFERENCE: P1661R2
CURRENT FILING DATE: US/10/614,959
CURRENT FILING DATE: 2003-07-08
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/383,667
PRIOR FILING DATE: EARLIER FILING DATE: 1999-08-26
PRIOR PHILING DATE: EARLIER FILING DATE: 1998-08-28
PRIOR PHILING DATE: EARLIER FILING DATE: 1998-08-03
PRIOR PHILING DATE: EARLIER FILING DATE: 1998-08-03
PRIOR PHILING DATE: EARLIER FILING DATE: 1998-03-03
PRIOR PHILING DATE: EARLIER FILING DATE: 1999-03-03
 CURRENT APPLICATION NUMBER: US/11/038,754
CURRENT FILING DATE: 2005-01-19
 NUMBER OF SEQ ID NOS: 32
 TYPE: PRT
 y Match 27.9%;
Local Similarity 50.0%;
hes 3; Conservative
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 5 LGEEYV 10
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 4 YLGEEYVK 11
 IGNNYV 12
 YLOREFEK 13
 Application US/10614959 to. US20060088534A1
 Conservative
 29.4%;
 1; Mismatches
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 Score 19; DB 6;
Pred. No. 9.2e+02;
 Score 20; [
Pred. No. 6.
 DB 7;
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 Length 13;
 2; Indels
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 Gaps
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 0
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APPLICANT: Epinmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOSTWARE: Patentin version 3.2
SEQ ID NO 35
LENGTH: 13
 APPLICANT: Lee, Liana
APPLICANT: Krystek, Stanley
APPLICANT: Krystek, Stanley
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A 1
TITLE OF INVENTION: IN THE TESTIS, MMP-29
FILE REFERENCE: DO141DIV
CURRENT APPLICATION NUMBER: US/11/313,356
CURRENT APPLICATION NUMBER: US/11/313,356
CURRENT FILING DATE: 2005-12-21
PRIOR APPLICATION NUMBER: US 60/286,764
PRIOR FILING DATE: 2001-04-26
NUMBER OF SEQ ID NOS: 80
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 S
 US-10-538-066-35
 RESULT 10
 ; ORGANISM: Homo sapiens
US-11-313-356-20
 US-11-264-509A-94
 Sequence 35, Application US/10538066 Publication No. US20060094649A1
 SOFTWARE: PatentIn version 3.2
SEQ ID NO 20
LENGTH: 13
TYPE: PRT
 APPLICANT: Wu, Shujian
APPLICANT: Chen, Jian
APPLICANT: Feder, John
APPLICANT: Lee, Liana
APPLICANT: Krystek, Star
 NUMBER OF SEQ ID NOS: 141
SOFTWARE: PatentIn version 3.3
SEQ ID NO 94
 GENERAL INFORMATION:
 GENERAL INFORMATION:
 Sequence 20, Application US/11313356
Publication No. US20060105379A1
 Matches
 Query Match
Best Local :
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 Best Local Similarity
 Query Match
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: PRT
TYPE: PRT
ORGANISM: Unknown
 OTHER INFORMATION: HCV peptide
 ENGTH:
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 Similarity 2; Conserv
 4; Conservative
 KKFISEKW 9
 EKYLGEEY 9
 EYVKAV 13
 EYVLAL 10
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 27.9%;
 27.9%;
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Pred. No. 9.
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 DB 7;
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 A NOVEL METALLOPROTEASE HIGHLY EXPRESSED
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 Length 13;
 and -B44
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 Associated
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FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
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LENGTH: 13
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US-10-538-066-32
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 ; OTHER INFORMATION: PADRE Peptide US-10-538-066-35
 RESULT 12
US-10-538-066-651
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 S
 밁
 ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-538-066-651
 ; OTHER INFORMATION: PADRE Peptide US-10-538-066-32
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Matches 3
 Sequence 651, Application US/10538066 Publication No. US20060094649A1 GENERAL INFORMATION:
 Sequence 32, Application US/10538066
Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and -B44 Tumor Associated Antigen
TITLE OF INVENTION: Peptides and Compositions
 Query Match
Best Local Similarity
Matches 4; Conserv
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 651
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 APPLICANT: Epinmune Inc.
APPLICANT: Epinmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.0159C06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR APPLICATION NUMBER: US 60/432,017
NUMBER OF SEQ ID NOS: 767
NUMBER OF SEQ ID NOS: 767
 TYPE: PRT
ORGANISM: Unknown
 FEATURE:
 LENGTH: 13
 Local Similarity 40.0
 N
 N
 3 KYLGEEYVKA 12
 8 EYVKAV 13
 ω
••
 Similarity
 EYLQLV 7
 KYLGEEYVKA 12
 KYVAAWTLKA 11
 KYVAAYTLKA 11
 Conservative
 Conservative
 25.0%;
 26.5%;
 25.0%;
 Score 18; DB 6;
Pred. No. 1.4e+03;
2; Mismatches 4
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 <u>ب</u>
 Score 17; DB 6;
Pred. No. 2e+03;
2; Mismatches
 Score 17; DB 6;
Pred. No. 2e+03;
2; Mismatches
 ŗ
 Length 13;
 Length 13;
 Length 13;
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 Indels
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FILE REFERENCE: JB06227US01
CURRENT APPLICATION NUMBER: US/11/264,509A
CURRENT FILING DATE: 2005-11-01
PRIOR APPLICATION NUMBER: 60/624,204
PRIOR FILING DATE: 2004-11-02
PRIOR FILING DATE: 2004-11-02
NUMBER OF SEQ ID NOS: 141
SOFTWARE: PatentIn version 3.3
SEQ ID NO 138
LENGTH: 13
SEC ID NO 138
LENGTH: 13
RESULT 15
US-11-261-429-21
US-11-261-429-21
Sequence 21, Application US/11261429
Publication No. US20060115899A1
GENERAL INFORMATION:
 Query Match
Best Local Similarity
Thehes 3; Conserve
 Query Match
Best Local Similarity
Matches 3; Conserv:
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 5
 ; OTHER INFORMATION: HCV peptide US-11-264-509A-128
 ; Publication No. US20060
; GENERAL INFORMATION:
; APPLICANT: Liu, Rong
; APPLICANT: Zhang, Rum
; APPLICANT: Kong, Rong
 RESULT 13
US-11-264-509A-128
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 US-11-264-509A-138
 US-11-264-509A-138
 RESULT 14
 Sequence 138, Application US/11264509A Publication No. US20060100148A1 GENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/11/264,509A
CURRENT FILING DATE: 2005-11-01
PRIOR APPLICATION NUMBER: 60/624,204
PRIOR FILING DATE: 2004-11-02
NUMBER OF SEQ ID NOS: 141
SOFTWARE: Patentin version 3.3
SEQ ID NO 128
LENGTH: 13
 Sequence 128, Application US/11264509A Publication No. US20060100148A1
 APPLICANT: Zhang, Rumin
APPLICANT: Kong, Rong
TITLE OF INVENTION: PEPTIDE VIRAL ENTRY PEPTIDES
 APPLICANT: Liu, Rong
 FILE REFERENCE: JB06227US01
 TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: HCV peptide
 ORGANISM: Artificial Sequence
 PPLICANT: Zhang, Rumin
PPLICANT: Kong, Rong
ITLE OF INVENTION: PEPTIDE VIRAL ENTRY PEPTIDES
 1 YEKYL 5
 7 EEYV 10
 FEVYL 8
 Conservative
 Conservative
 Rumin
 25.0%;
 25.0%;
 1; Mismatches
 Score 17; DB 7;
Pred. No. 2e+03;
 Score 17; DB 7;
Pred. No. 2e+03;
 Mismatches
 DB 7;
 Length 13;
 Length 13
 Indels
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
```

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APPLICANT: Walker, Mindi R.

TITILE OF INVENTION: METHODS OF GENERATING ANTIGEN-SPECIFIC CD4+CD25+ REGULATORY T

TITLE OF INVENTION: CELLS, COMPOSITIONS AND METHODS OF USE

FILE REFERENCE: BRIVM-1-26413

CURRENT APPLICATION NUMBER: US/1/261,429

CURRENT APPLICATION NUMBER: US/210-21

PRIOR APPLICATION NUMBER: US/60/623,380

PRIOR FILLING DATE: 2005-10-28

NUMBER OF SEQ ID NOS: 65

SOFTWARE: PATCHIT VETSION 3.2

SEQ ID NO 21

LENGTH: 13

TYPE: PRT

ORGANISM: Homo Sapiens

US-11-261-429-21

Query Match

Best Local Similarity 75.0%; Score 17; DB 7; Length 13;

Matches 3; COnservative 1; Mismatches 0; Indels 0; Gaps 0;

Matches 3; COnservative 1; Mismatches 0; Indels 0; Gaps 0;

Search completed: June 13, 2006, 03:35:20

Job time: 38 secs
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